



SEQUENCE LISTING

<110> DEHESH et al.

<120> Engineering Beta Ketoacyl ACP Synthase for Novel Substrate Specificity

<130> 16516.117

<140> US 09/591,279

<141> 2000-06-09

<150> US 60/138,308

<151> 1999-06-09

<160> 46

<170> PatentIn version 3.0

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<213> Artificial Sequence

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<223> Oligonucleotide Primer I108F Sense

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36

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36

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<211> 42

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42

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cgatcagtcc gaggcctcca agcccggagc caattgcggc ac

42

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gcaggtggcg ccgagaaaat cagtacgccg ctgggc

36

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gcccagcggc gtactgattt tctcggcgcc acctg

35

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ggtggcgag agaaaatgag tactccgctg ggcgttg 37

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caacgcccag cggagtactc attttctctg cgccacc 37

<210> 9
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<223> Oligonucleotide Primer I108A,L111A, I114A Sense

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gcaattggct cgggggctgg cggcgccgga ctggccgaag aaaaccacac 50

<210> 10
<211> 50
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<223> Oligonucleotide Primer L111A Sense

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gggattggcg ggcgcggact gatcgaag

28

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cttcgatcag tccggcgccg ccaatccc

28

<210> 13

<211> 34

<212> DNA

<213> Artificial Sequence

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gacgagccca ttcgcggtac cgtcaacgat tgtg

34

<210> 14

<211> 34

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cacaatcggt gacggtaccg cgaatgggct gatc

34

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 <211> 32
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<400> 15
 gagaaagcca gtactccggc gggcgttggt gg 32

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<210> 17
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<220>
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 <223> Self annealed oligonucleotide primer

<400> 17
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<210> 18
 <211> 366
 <212> DNA
 <213> Cuphea hookeriana

<400> 18
 ctgagatctg tcgacatggc gaccgcttct cgcattggtg cgtccccctt ctgtacgtgg 60

ctcgtagctg catgcatgcc cacttcatcc gacaacgacc cagttccct tccccacaag 120

cggtccggc tetccgctg ccggaggact ctctctccc attgctccct ccgcggtacc 180

accttccaat gcctcgatcc ttgcaaccag caacgcttcc tcggggataa cggattoct 240
 tccctcttcg gatccaagcc tcttcgttca aatcgcggcc acctgaggct cggccgcact 300
 tcccattccg gggagggtcat ggctgtggct atgcaacctg cacaggaagt ctccacaaga 360
 tctgtc 366

<210> 19
 <211> 431
 <212> PRT
 <213> Arabidopsis thaliana

<400> 19

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Pro	Lys	Lys	Arg	Val	Val	Ile	Thr	Gly	Met	Gly	Leu	Val	Ser	Val	Cys	20	25	30	
Gly	Asn	Asp	Val	Asp	Ala	Tyr	Tyr	Glu	Lys	Leu	Leu	Ser	Gly	Glu	Ser	35	40	45	
Gly	Ile	Ser	Leu	Ile	Asp	Arg	Phe	Asp	Ala	Ser	Lys	Phe	Pro	Thr	Arg	50	55	60	
Phe	Gly	Gly	Gln	Ile	Arg	Gly	Phe	Ser	Ser	Glu	Gly	Tyr	Ile	Asp	Gly	65	70	75	80
Lys	Asn	Glu	Arg	Arg	Leu	Asp	Asp	Cys	Leu	Lys	Tyr	Cys	Ile	Val	Ala	85	90	95	
Gly	Lys	Lys	Ala	Leu	Glu	Ser	Ala	Asn	Leu	Gly	Gly	Asp	Lys	Leu	Asn	100	105	110	
Thr	Ile	Asp	Lys	Arg	Lys	Ala	Gly	Val	Leu	Val	Gly	Thr	Gly	Met	Gly	115	120	125	
Gly	Leu	Thr	Val	Phe	Ser	Glu	Gly	Val	Gln	Asn	Leu	Ile	Glu	Lys	Gly	130	135	140	
His	Arg	Arg	Ile	Ser	Pro	Phe	Phe	Ile	Pro	Tyr	Ala	Ile	Thr	Asn	Met	145	150	155	160
Gly	Ser	Ala	Leu	Leu	Ala	Ile	Asp	Leu	Gly	Leu	Met	Gly	Pro	Asn	Tyr	165	170	175	
Ser	Ile	Ser	Thr	Ala	Cys	Ala	Thr	Ser	Asn	Tyr	Cys	Phe	Tyr	Ala	Ala	180	185	190	
Ala	Asn	His	Asn	His	Arg	Gly	Glu	Ala	Asp	Met	Met	Ile	Ala	Gly	Gly	195	200	205	
Thr	Glu	Ala	Ala	Ile	Ile	Pro	Ile	Gly	Leu	Gly	Gly	Phe	Val	Ala	Cys	210	215	220	

Arg Ala Leu Ser Gln Arg Asn Asp Asp Pro Gln Thr Ala Ser Arg Pro
 225 230 235 240
 Trp Asp Lys Ala Arg Asp Gly Phe Val Met Gly Glu Gly Ala Gly Val
 245 250 255
 Leu Val Met Glu Ser Leu Glu His Ala Met Lys Arg Gly Ala Pro Ile
 260 265 270
 Val Ala Glu Tyr Leu Gly Gly Ala Val Asn Cys Asp Ala His His Met
 275 280 285
 Thr Asp Pro Arg Ala Asp Gly Leu Gly Val Ser Ser Cys Ile Glu Arg
 290 295 300
 Cys Leu Glu Asp Ala Gly Val Ser Pro Glu Glu Val Asn Tyr Ile Asn
 305 310 315 320
 Ala His Ala Thr Ser Thr Leu Ala Gly Asp Leu Ala Glu Ile Asn Ala
 325 330 335
 Ile Lys Lys Val Phe Lys Ser Thr Ser Gly Ile Lys Ile Asn Ala Thr
 340 345 350
 Lys Ser Met Ile Gly His Cys Leu Gly Ala Ala Gly Gly Leu Glu Ala
 355 360 365
 Ile Ala Thr Val Lys Ala Ile Asn Thr Gly Trp Leu His Pro Ser Ile
 370 375 380
 Asn Gln Phe Asn Pro Glu Gln Ala Val Asp Phe Asp Thr Val Pro Asn
 385 390 395 400
 Glu Lys Lys Gln His Glu Val Asp Val Ala Ile Ser Asn Ser Phe Gly
 405 410 415
 Phe Gly Gly His Asn Ser Val Val Ala Phe Ser Ala Phe Lys Pro
 420 425 430

<210> 20
 <211> 429
 <212> PRT
 <213> Brassica napus

<400> 20

Ala Ser Ser Ser Ala Val Ser Ala Pro Lys Arg Glu Thr Asp Pro Lys
 1 5 10 15
 Lys Arg Val Val Ile Thr Gly Met Gly Leu Val Ser Val Phe Gly Asn
 20 25 30
 Asp Val Asp Ala Tyr Tyr Glu Lys Leu Leu Ser Gly Glu Ser Gly Ile
 35 40 45
 Ser Leu Ile Asp Arg Phe Asp Ala Ser Lys Phe Pro Thr Arg Phe Gly

50					55					60						
Gly	Gln	Ile	Arg	Gly	Phe	Ser	Ser	Glu	Gly	Tyr	Ile	Asp	Gly	Lys	Asn	
65					70					75					80	
Glu	Arg	Arg	Leu	Asp	Asp	Cys	Leu	Lys	Tyr	Cys	Ile	Val	Ala	Gly	Lys	
				85					90					95		
Lys	Ala	Leu	Glu	Ser	Ala	Asn	Leu	Gly	Gly	Asp	Lys	Leu	Asn	Thr	Ile	
			100					105					110			
Asp	Lys	Gln	Lys	Ala	Gly	Val	Leu	Val	Gly	Thr	Gly	Met	Gly	Gly	Leu	
		115					120					125				
Thr	Val	Phe	Ser	Asp	Gly	Val	Gln	Ala	Leu	Ile	Glu	Lys	Gly	His	Arg	
		130					135					140				
Arg	Ile	Ser	Pro	Phe	Phe	Ile	Pro	Tyr	Ala	Ile	Thr	Asn	Met	Gly	Ser	
				150								155			160	
Ala	Leu	Leu	Ala	Ile	Asp	Leu	Gly	Leu	Met	Gly	Pro	Asn	Tyr	Ser	Ile	
				165					170					175		
Ser	Thr	Ala	Cys	Ala	Thr	Ser	Asn	Tyr	Cys	Phe	Tyr	Ala	Ala	Ala	Asn	
			180					185					190			
His	Ile	Arg	Arg	Gly	Glu	Ala	Asp	Met	Met	Ile	Ala	Gly	Gly	Thr	Glu	
		195					200					205				
Ala	Ala	Ile	Ile	Pro	Ile	Gly	Leu	Gly	Gly	Phe	Val	Ala	Cys	Arg	Ala	
		210					215					220				
Leu	Ser	Gln	Arg	Asn	Asp	Asp	Pro	Gln	Thr	Ala	Ser	Arg	Pro	Trp	Asp	
				230								235			240	
Lys	Gln	Arg	Asp	Gly	Phe	Val	Met	Gly	Glu	Gly	Ala	Gly	Val	Leu	Val	
				245					250					255		
Met	Glu	Ser	Leu	Glu	His	Ala	Met	Lys	Arg	Gly	Ala	Pro	Ile	Val	Ala	
			260					265					270			
Glu	Tyr	Leu	Gly	Gly	Ala	Val	Asn	Cys	Asp	Ala	His	His	Met	Thr	Asp	
		275					280					285				
Pro	Arg	Ala	Asp	Gly	Leu	Gly	Val	Ser	Ser	Cys	Ile	Glu	Ser	Cys	Leu	
				290			295					300				
Glu	Asp	Ala	Gly	Val	Ser	Pro	Glu	Glu	Val	Asn	Tyr	Ile	Asn	Ala	His	
				310								315			320	
Ala	Thr	Ser	Thr	Leu	Ala	Gly	Asp	Leu	Ala	Glu	Ile	Asn	Ala	Ile	Lys	
				325					330					335		
Lys	Val	Phe	Lys	Ser	Thr	Ser	Gly	Ile	Lys	Ile	Asn	Ala	Thr	Lys	Ser	
			340					345					350			
Met	Ile	Gly	His	Cys	Leu	Gly	Ala	Ala	Gly	Gly	Leu	Glu	Ala	Ile	Ala	

355	360	365
Thr Val Lys Ala Ile Asn Thr Gly Trp Leu His Pro Ser Ile Asn Gln 370 375 380		
Phe Asn Pro Glu Pro Ala Val Asp Phe Asp Thr Val Ala Asn Glu Lys 385 390 395 400		
Lys Gln His Glu Val Asn Val Ala Ile Ser Asn Ser Phe Gly Phe Gly 405 410 415		
Gly His Asn Ser Val Val Ala Phe Ser Ala Phe Lys Pro 420 425		
<210> 21		
<211> 350		
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<213> Cuphea hookeriana		
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Ser Ser Thr Ala Val Ala Ala Ala Leu Glu Leu Val Asp Pro Pro Gly 1 5 10 15		
Cys Arg Asn Ser Ala Arg Ala Asp Leu Gly Ala Asp Arg Leu Ser Lys 20 25 30		
Ile Asp Lys Glu Arg Ala Gly Val Leu Val Gly Thr Gly Met Gly Gly 35 40 45		
Leu Thr Val Phe Ser Asp Gly Val Gln Ser Leu Ile Glu Lys Gly His 50 55 60		
Arg Lys Ile Thr Pro Phe Phe Ile Pro Tyr Ala Ile Thr Asn Met Gly 65 70 75 80		
Ser Ala Leu Leu Ala Ile Glu Phe Gly Leu Met Gly Pro Asn Tyr Ser 85 90 95		
Ile Ser Thr Ala Cys Ala Thr Ser Asn Tyr Cys Phe His Ala Ala Ala 100 105 110		
Asn His Ile Arg Arg Gly Glu Ala Asp Leu Met Ile Ala Gly Gly Thr 115 120 125		
Glu Ala Ala Ile Ile Pro Ile Gly Leu Gly Gly Phe Val Ala Cys Arg 130 135 140		
Ala Leu Ser Gln Arg Asn Asp Asp Pro Gln Thr Ala Ser Arg Pro Trp 145 150 155 160		
Asp Lys Asp Arg Asp Gly Phe Val Met Gly Glu Gly Ala Gly Val Leu 165 170 175		
Val Met Glu Ser Leu Glu His Ala Met Arg Arg Gly Ala Pro Ile Ile 180 185 190		

Ala Glu Tyr Leu Gly Gly Ala Ile Asn Cys Asp Ala Tyr His Met Thr
195 200 205

Asp Pro Arg Ala Asp Gly Leu Gly Val Ser Ser Cys Ile Glu Ser Ser
210 215 220

Leu Glu Asp Ala Gly Val Ser Pro Glu Glu Val Asn Tyr Ile Asn Ala
225 230 235 240

His Ala Thr Ser Thr Leu Ala Gly Asp Leu Ala Glu Ile Asn Ala Ile
245 250 255

Lys Lys Val Phe Lys Asn Thr Lys Asp Ile Lys Ile Asn Ala Thr Lys
260 265 270

Ser Met Ile Gly His Cys Leu Gly Ala Ser Gly Gly Leu Glu Ala Ile
275 280 285

Ala Thr Ile Lys Gly Ile Asn Thr Gly Trp Leu His Pro Ser Ile Asn
290 295 300

Gln Phe Asn Pro Glu Pro Ser Val Glu Phe Asp Thr Val Ala Asn Lys
305 310 315 320

Lys Gln Gln His Glu Val Asn Val Ala Ile Ser Asn Ser Phe Gly Phe
325 330 335

Gly Gly His Asn Ser Val Val Ala Phe Ser Ala Phe Lys Pro
340 345 350

<210> 22
<211> 441
<212> PRT
<213> Cuphea hookeriana

<220>
<221> misc_feature
<222> (15)..(15)
<223> Xaa at position 15 is unknown.

<400> 22

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Ala Ala Ala Leu Glu Leu Val Asp Pro Pro Gly Cys Arg Asn Ser Ala
20 25 30

Arg Ala Gly Met Gly Leu Val Ser Val Phe Gly Ser Asp Val Asp Ser
35 40 45

Tyr Tyr Glu Lys Leu Leu Ser Gly Glu Ser Gly Ile Ser Leu Ile Asp
50 55 60

Arg Phe Asp Ala Ser Lys Phe Pro Thr Arg Phe Gly Gly Gln Ile Arg
65 70 75 80

Gly	Phe	Asn	Ala	Thr	Gly	Tyr	Ile	Asp	Gly	Lys	Asn	Asp	Arg	Arg	Leu	85	90	95
Asp	Asp	Cys	Leu	Arg	Tyr	Cys	Ile	Val	Ala	Gly	Lys	Lys	Ala	Leu	Glu	100	105	110
Asn	Ser	Asp	Leu	Gly	Gly	Glu	Ser	Leu	Ser	Lys	Ile	Asp	Lys	Glu	Arg	115	120	125
Ala	Gly	Val	Leu	Val	Gly	Thr	Gly	Met	Gly	Gly	Leu	Thr	Val	Phe	Ser	130	135	140
Asp	Gly	Val	Gln	Asn	Leu	Ile	Glu	Lys	Gly	His	Arg	Lys	Ile	Ser	Pro	145	150	155
Phe	Phe	Ile	Pro	Tyr	Ala	Ile	Thr	Asn	Met	Gly	Ser	Ala	Leu	Leu	Ala	165	170	175
Ile	Asp	Leu	Gly	Leu	Met	Gly	Pro	Asn	Tyr	Ser	Ile	Ser	Thr	Ala	Cys	180	185	190
Ala	Thr	Ser	Asn	Tyr	Cys	Phe	Tyr	Ala	Ala	Ala	Asn	His	Ile	Arg	Arg	195	200	205
Gly	Glu	Ala	Asp	Leu	Met	Ile	Ala	Gly	Gly	Thr	Glu	Ala	Ala	Ile	Ile	210	215	220
Pro	Ile	Gly	Leu	Gly	Gly	Phe	Val	Ala	Cys	Arg	Ala	Leu	Ser	Gln	Arg	225	230	235
Asn	Asp	Asp	Pro	Gln	Thr	Ala	Ser	Arg	Pro	Trp	Asp	Lys	Asp	Arg	Asp	245	250	255
Gly	Phe	Val	Met	Gly	Glu	Gly	Ala	Gly	Val	Leu	Val	Met	Glu	Ser	Leu	260	265	270
Glu	His	Ala	Met	Lys	Arg	Gly	Ala	Pro	Ile	Ile	Ala	Glu	Tyr	Leu	Gly	275	280	285
Gly	Ala	Val	Asn	Cys	Asp	Ala	Tyr	His	Met	Thr	Asp	Pro	Arg	Ala	Asp	290	295	300
Gly	Leu	Gly	Val	Ser	Ser	Cys	Ile	Glu	Ser	Ser	Leu	Glu	Asp	Ala	Gly	305	310	315
Val	Ser	Pro	Glu	Glu	Val	Asn	Tyr	Ile	Asn	Ala	His	Ala	Thr	Ser	Thr	325	330	335
Leu	Ala	Gly	Asp	Leu	Ala	Glu	Ile	Asn	Ala	Ile	Lys	Lys	Val	Phe	Lys	340	345	350
Asn	Thr	Lys	Glu	Ile	Thr	Ile	Asn	Ala	Thr	Lys	Ser	Met	Ile	Gly	His	355	360	365
Cys	Leu	Gly	Ala	Ser	Gly	Gly	Leu	Glu	Ala	Ile	Ala	Thr	Ile	Lys	Gly	370	375	380

Ile Thr Thr Gly Trp Leu His Pro Ser Ile Asn Gln Phe Asn Pro Glu
 385 390 395 400

Pro Ser Val Glu Phe Asp Thr Val Ala Asn Lys Lys Gln Gln His Glu
 405 410 415

Val Asn Val Ala Ile Ser Asn Ser Phe Gly Phe Gly Gly His Asn Ser
 420 425 430

Val Val Ala Phe Ser Ala Phe Lys Pro
 435 440

<210> 23
 <211> 430
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 <213> Cuphea pullcherima

<400> 23

Arg Ala Ala Ser Pro Thr Val Ser Ala Pro Lys Arg Glu Thr Asp Pro
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Lys Lys Arg Val Val Ile Thr Gly Met Gly Leu Val Ser Val Phe Gly
 20 25 30

Ser Asp Val Asp Ala Tyr Tyr Asp Lys Leu Leu Ser Gly Glu Ser Gly
 35 40 45

Ile Gly Pro Ile Asp Arg Phe Asp Ala Ser Lys Phe Pro Thr Arg Phe
 50 55 60

Gly Gly Gln Ile Arg Gly Phe Asn Ser Met Gly Tyr Ile Asp Gly Lys
 65 70 75 80

Asn Asp Arg Arg Leu Asp Asp Cys Leu Arg Tyr Cys Ile Val Ala Gly
 85 90 95

Lys Lys Ser Leu Glu Asp Ala Asp Leu Gly Ala Asp Arg Leu Ser Lys
 100 105 110

Ile Asp Lys Glu Arg Ala Gly Val Leu Val Gly Thr Gly Met Gly Gly
 115 120 125

Leu Thr Val Phe Ser Asp Gly Val Gln Ser Leu Ile Glu Lys Gly His
 130 135 140

Arg Lys Ile Thr Pro Phe Phe Ile Pro Tyr Ala Ile Thr Asn Met Gly
 145 150 155 160

Ser Ala Leu Leu Ala Ile Glu Leu Gly Leu Met Gly Pro Asn Tyr Ser
 165 170 175

Ile Ser Thr Ala Cys Ala Thr Ser Asn Tyr Cys Phe His Ala Ala Ala
 180 185 190

Asn His Ile Arg Arg Gly Glu Ala Asp Leu Met Ile Ala Gly Gly Thr

195					200					205					
Glu	Ala	Ala	Ile	Ile	Pro	Ile	Gly	Leu	Gly	Gly	Phe	Val	Ala	Cys	Arg
210						215					220				
Ala	Leu	Ser	Gln	Arg	Asn	Asp	Asp	Pro	Gln	Thr	Ala	Ser	Arg	Pro	Trp
225					230					235					240
Asp	Lys	Asp	Arg	Asp	Gly	Phe	Val	Met	Gly	Glu	Gly	Ala	Gly	Val	Leu
				245					250					255	
Val	Leu	Glu	Ser	Leu	Glu	His	Ala	Met	Lys	Arg	Gly	Ala	Pro	Ile	Ile
			260					265					270		
Ala	Glu	Tyr	Leu	Gly	Gly	Ala	Ile	Asn	Cys	Asp	Ala	Tyr	His	Met	Thr
	275						280					285			
Asp	Pro	Arg	Ala	Asp	Gly	Leu	Gly	Val	Ser	Ser	Cys	Ile	Glu	Ser	Ser
	290					295					300				
Leu	Glu	Asp	Ala	Gly	Val	Ser	Pro	Glu	Glu	Val	Asn	Tyr	Ile	Asn	Ala
305					310					315					320
His	Ala	Thr	Ser	Thr	Leu	Ala	Gly	Asp	Leu	Ala	Glu	Ile	Asn	Ala	Ile
				325					330					335	
Lys	Lys	Val	Phe	Lys	Asn	Thr	Lys	Asp	Ile	Lys	Ile	Asn	Ala	Thr	Lys
			340					345					350		
Ser	Met	Ile	Gly	His	Cys	Leu	Gly	Ala	Ser	Gly	Gly	Leu	Glu	Ala	Ile
	355						360					365			
Ala	Thr	Ile	Lys	Gly	Ile	Asn	Thr	Gly	Trp	Leu	His	Pro	Ser	Ile	Asn
	370					375					380				
Gln	Phe	Asn	Pro	Glu	Pro	Ser	Val	Glu	Phe	Asp	Thr	Val	Ala	Asn	Lys
385					390					395					400
Lys	Gln	Gln	His	Glu	Val	Asn	Val	Ala	Ile	Ser	Asn	Ser	Phe	Gly	Phe
				405					410					415	
Gly	Gly	His	Asn	Ser	Val	Val	Ala	Phe	Ser	Ala	Phe	Lys	Pro		
			420					425					430		

<210> 24
 <211> 428
 <212> PRT
 <213> Cuphea pullcherima

<400> 24

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Arg	Val	Val	Ile	Thr	Gly	Met	Gly	Leu	Val	Ser	Val	Phe	Gly	Ser	Asp
			20					25					30		

Val	Asp	Ala	Tyr	Tyr	Asp	Lys	Leu	Leu	Ser	Gly	Glu	Ser	Gly	Ile	Ser		
	35						40					45					
Leu	Ile	Asp	Arg	Phe	Asp	Ala	Ser	Lys	Phe	Pro	Thr	Arg	Phe	Ala	Gly		
	50					55					60						
Gln	Ile	Arg	Gly	Phe	Asn	Ala	Thr	Gly	Tyr	Ile	Asp	Gly	Lys	Asn	Asp		
	65				70				75					80			
Arg	Arg	Leu	Asp	Asp	Cys	Leu	Arg	Tyr	Cys	Ile	Val	Ala	Gly	Lys	Lys		
			85					90						95			
Ala	Leu	Glu	Asp	Ala	Asp	Leu	Ala	Gly	Gln	Ser	Leu	Ser	Lys	Ile	Asp		
			100					105					110				
Lys	Glu	Arg	Ala	Gly	Val	Leu	Val	Gly	Thr	Gly	Met	Gly	Gly	Leu	Thr		
	115					120						125					
Val	Phe	Ser	Asp	Gly	Val	Gln	Asn	Leu	Ile	Glu	Lys	Gly	His	Arg	Lys		
	130					135					140						
Ile	Ser	Pro	Phe	Phe	Ile	Pro	Tyr	Ala	Ile	Thr	Asn	Met	Gly	Ser	Ala		
	145				150					155					160		
Leu	Leu	Ala	Ile	Asp	Leu	Gly	Leu	Met	Gly	Pro	Asn	Tyr	Ser	Ile	Ser		
			165					170						175			
Thr	Ala	Cys	Ala	Thr	Ser	Asn	Tyr	Cys	Phe	Tyr	Ala	Ala	Ala	Asn	His		
		180						185					190				
Ile	Arg	Arg	Gly	Glu	Ala	Asp	Leu	Met	Ile	Ala	Gly	Gly	Thr	Glu	Ala		
	195					200						205					
Ala	Val	Ile	Pro	Ile	Gly	Leu	Gly	Gly	Phe	Val	Ala	Cys	Arg	Ala	Leu		
	210					215					220						
Ser	Gln	Arg	Asn	Asp	Asp	Pro	Gln	Thr	Ala	Ser	Arg	Pro	Trp	Asp	Lys		
	225			230						235				240			
Asp	Arg	Asp	Gly	Phe	Val	Met	Gly	Glu	Gly	Ala	Gly	Val	Leu	Val	Met		
			245					250					255				
Glu	Ser	Leu	Glu	His	Ala	Met	Lys	Arg	Gly	Ala	Pro	Ile	Ile	Ala	Glu		
		260						265					270				
Tyr	Leu	Gly	Gly	Ala	Val	Asn	Cys	Asp	Ala	Tyr	His	Met	Thr	Asp	Pro		
	275					280						285					
Arg	Ala	Asp	Gly	Leu	Gly	Val	Ser	Ser	Cys	Ile	Glu	Ser	Ser	Leu	Glu		
	290					295					300						
Asp	Ala	Gly	Val	Ser	Pro	Glu	Glu	Val	Asn	Tyr	Ile	Asn	Ala	His	Ala		
	305				310					315					320		
Thr	Ser	Thr	Leu	Ala	Gly	Asp	Leu	Ala	Glu	Ile	Asn	Ala	Ile	Lys	Lys		
			325					330						335			

Val Phe Lys Asn Thr Lys Glu Ile Lys Ile Asn Ala Thr Lys Ser Met
 340 345 350

Ile Gly His Cys Leu Gly Ala Ser Gly Gly Leu Glu Ala Ile Ala Thr
 355 360 365

Ile Lys Gly Ile Thr Thr Gly Trp Leu His Pro Ser Ile Asn Gln Phe
 370 375 380

Asn Pro Glu Pro Ser Val Asp Phe Asn Thr Val Ala Asn Lys Lys Gln
 385 390 395 400

Gln His Glu Val Asn Val Ala Ile Ser Asn Ser Phe Gly Phe Gly Gly
 405 410 415

His Asn Ser Val Val Ala Phe Ser Ala Phe Lys Pro
 420 425

<210> 25
 <211> 427
 <212> PRT
 <213> Hordeum vulgare

<400> 25

Thr Ser Ala Ala Pro Gln Arg Glu Thr Asp Pro Arg Lys Arg Val Val
 1 5 10 15

Ile Thr Gly Met Gly Leu Ala Ser Val Phe Gly Ser Asp Val Asp Thr
 20 25 30

Phe Tyr Asp Arg Leu Leu Ala Gly Glu Ser Gly Val Gly Pro Ile Asp
 35 40 45

Arg Phe Asp Ala Ser Ser Phe Pro Thr Arg Phe Ala Gly Gln Ile Arg
 50 55 60

Gly Phe Ser Ser Glu Gly Tyr Ile Asp Gly Lys Asn Asp Arg Arg Leu
 65 70 75 80

Asp Asp Cys Ile Arg Tyr Cys Ile Leu Ser Gly Lys Lys Ala Leu Glu
 85 90 95

Ser Ala Gly Leu Gly Ala Gly Ser Asp Ala His Val Lys Leu Asp Val
 100 105 110

Gly Arg Ala Gly Val Leu Val Gly Thr Gly Met Gly Gly Leu Ser Val
 115 120 125

Phe Ser Asp Gly Val Gln Asn Leu Ile Glu Lys Gly Tyr Arg Lys Ile
 130 135 140

Ser Pro Phe Phe Ile Pro Tyr Ala Ile Thr Asn Met Gly Ser Ala Leu
 145 150 155 160

Leu Ala Ile Asp Val Gly Phe Met Gly Pro Asn Tyr Ser Ile Ser Thr
 165 170 175

Ala Cys Ala Thr Ser Asn Tyr Cys Phe Tyr Ala Ala Ala Asn His Ile
180 185 190

Arg Arg Gly Glu Ala Asp Ile Ile Val Ala Gly Gly Thr Glu Ala Ala
195 200 205

Ile Ile Pro Ile Gly Leu Gly Gly Phe Val Ala Cys Arg Ala Leu Ser
210 215 220

Gln Arg Asn Asp Asp Pro Ile Thr Ala Cys Arg Pro Trp Asp Lys Glu
225 230 235 240

Arg Asp Gly Phe Val Met Gly Glu Gly Ala Gly Val Leu Val Met Glu
245 250 255

Ser Leu Glu His Ala Met Lys Arg Asp Ala Pro Ile Ile Ala Glu Tyr
260 265 270

Leu Gly Gly Ala Val Asn Cys Asp Ala Tyr His Met Thr Asp Pro Arg
275 280 285

Ala Asp Gly Leu Gly Val Ser Ser Cys Ile Thr Met Ser Leu Arg Asp
290 295 300

Ala Gly Val Ala Pro Glu Glu Val Asn Tyr Ile Asn Ala His Ala Thr
305 310 315 320

Ser Thr Leu Ala Gly Asp Leu Ala Glu Val Arg Ala Ile Lys Gln Val
325 330 335

Phe Lys Asn Pro Ser Glu Ile Lys Ile Asn Ser Thr Lys Ser Met Ile
340 345 350

Gly His Cys Leu Gly Ala Ala Gly Gly Leu Glu Ala Ile Ala Thr Ile
355 360 365

Lys Ser Ile Thr Thr Gly Trp Val His Pro Thr Ile Asn Gln Phe Asn
370 375 380

Pro Glu Pro Glu Val Asp Phe Asp Thr Val Ala Asn Glu Lys Lys Gln
385 390 395 400

His Glu Val Asn Val Gly Ile Ser Asn Ser Phe Gly Phe Gly Gly His
405 410 415

Asn Ser Val Val Val Phe Ala Pro Phe Lys Pro
420 425

<210> 26
<211> 428
<212> PRT
<213> Ricinus communis

<400> 26

Asn Asn Asn Thr Thr Ile Ser Ala Pro Lys Arg Glu Lys Asp Pro Arg

1	5	10	15
Lys Arg Val Val Ile Thr Gly Thr Gly Leu Val Ser Val Phe Gly Asn	20	25	30
Asp Val Asp Thr Tyr Tyr Asp Lys Leu Leu Ala Gly Glu Ser Gly Ile	35	40	45
Gly Leu Ile Asp Arg Phe Asp Ala Ser Lys Phe Pro Thr Arg Phe Gly	50	55	60
Gly Gln Ile Arg Gly Phe Asn Ser Gln Gly Tyr Ile Asp Gly Lys Asn	65	70	75
Asp Arg Arg Leu Asp Asp Cys Leu Arg Tyr Cys Ile Val Ala Gly Lys	85	90	95
Lys Ala Leu Glu His Ala Asp Leu Gly Gly Asp Lys Leu Ser Lys Ile	100	105	110
Asp Lys Glu Arg Ala Gly Val Leu Val Gly Thr Gly Met Gly Gly Leu	115	120	125
Thr Val Phe Ser Asp Gly Val Gln Ala Leu Ile Glu Lys Gly His Arg	130	135	140
Lys Ile Thr Pro Phe Phe Ile Pro Tyr Ala Ile Thr Asn Met Gly Ser	145	150	155
Ala Leu Leu Ala Ile Glu Leu Gly Leu Met Gly Pro Asn Tyr Ser Ile	165	170	175
Ser Thr Ala Cys Ala Thr Ser Asn Tyr Cys Phe Tyr Ala Ala Ala Asn	180	185	190
His Ile Arg Arg Gly Glu Ala Glu Leu Met Ile Ala Gly Gly Thr Glu	195	200	205
Ala Ala Ile Ile Pro Ile Gly Leu Gly Gly Phe Val Ala Cys Arg Ala	210	215	220
Leu Ser Gln Arg Asn Asp Asp Pro Gln Thr Ala Ser Arg Pro Trp Asp	225	230	235
Lys Asp Arg Asp Gly Phe Val Met Gly Glu Gly Ala Gly Val Leu Val	245	250	255
Met Glu Ser Leu Glu His Ala Met Lys Arg Gly Ala Pro Ile Ile Ala	260	265	270
Glu Tyr Leu Gly Gly Ala Val Asn Cys Asp Ala Tyr His Met Thr Asp	275	280	285
Pro Arg Ala Asp Gly Leu Gly Val Ser Ser Cys Ile Glu Arg Ser Leu	290	295	300
Glu Asp Ala Gly Val Ser Pro Glu Glu Val Asn Tyr Ile Asn Ala His			

305		310		315		320
Ala Thr Ser Thr	Leu Ala Gly Asp	Leu Ala Glu Ile	Asn Ala Ile Lys			
	325		330			335
Lys Val Phe Lys	Asn Thr Ser Asp	Ile Lys Ile Asn	Ala Thr Lys Ser			
	340		345			350
Met Ile Gly His	Cys Leu Gly Ala	Ala Gly Gly Leu	Glu Ala Ile Ala			
	355		360			365
Cys Val Lys Ala	Ile Thr Thr Gly	Trp Leu His Pro	Thr Ile Asn Gln			
	370		375			380
Phe Asn Pro Glu	Pro Ser Val Glu	Phe Asp Thr Val	Ala Asn Lys Lys			
	385		390			395
Gln Gln His Glu	Val Asn Val Ala	Ile Ser Asn Ser	Phe Gly Phe Gly			
	405		410			415
Gly His Asn Ser	Val Val Ala Phe	Ser Ala Phe Lys				
	420		425			
<210> 27						
<211> 420						
<212> PRT						
<213> Capsicum chinense						
<400> 27						
Lys Arg Glu Thr	Asp Pro Lys Lys	Arg Ile Val Ile	Thr Gly Met Gly			
1	5	10	15			
Leu Val Ser Val	Phe Gly Ser Asp	Ile Asp Asn Phe	Tyr Asn Lys Leu			
	20	25	30			
Leu Glu Gly Gln	Ser Gly Ile Ser	Leu Ile Asp Arg	Phe Asp Ala Ser			
	35	40	45			
Ser Tyr Thr Val	Arg Phe Ala Gly	Gln Ile Arg Asp	Phe Ser Ser Glu			
	50	55	60			
Gly Tyr Ile Asp	Gly Lys Asn Asp	Arg Arg Leu Asp	Asp Cys Trp Arg			
65	70	75	80			
Tyr Cys Leu Val	Ala Gly Lys Arg	Ala Leu Glu Asp	Ala Asn Leu Gly			
	85	90	95			
Gln Gln Val Leu	Asp Thr Met Asp	Lys Thr Arg Ile	Gly Val Leu Val			
	100	105	110			
Gly Ser Ser Met	Gly Gly Ser Lys	Val Phe Ala Asp	Ala Val Glu Ala			
	115	120	125			
Leu Val Gln Arg	Gly Tyr Lys Lys	Ile Asn Pro Phe	Phe Ile Pro Tyr			
	130	135	140			

Ser	Ile	Thr	Asn	Met	Gly	Ser	Ala	Leu	Leu	Ala	Ile	Asp	Thr	Gly	Leu	145	150	155	160
Met	Gly	Pro	Thr	Tyr	Ser	Ile	Ser	Thr	Ala	Cys	Ala	Thr	Ala	Asn	Tyr	165	170	175	
Cys	Phe	Tyr	Ala	Ser	Ala	Asn	His	Ile	Arg	Arg	Gly	Glu	Ala	Asp	Ile	180	185	190	
Met	Val	Ala	Gly	Gly	Thr	Asp	Ala	Phe	Ile	Ser	Ala	Ile	Gly	Val	Gly	195	200	205	
Gly	Leu	Ile	Ala	Cys	Arg	Ala	Leu	Ser	Gln	Arg	Asn	Asp	Glu	Tyr	Glu	210	215	220	
Lys	Ala	Ser	Arg	Pro	Trp	Asp	Arg	Asn	Arg	Asp	Gly	Phe	Val	Ile	Gly	225	230	235	240
Glu	Gly	Ser	Gly	Val	Leu	Val	Met	Glu	Asn	Leu	Glu	His	Ala	Leu	Lys	245	250	255	
Arg	Gly	Ala	Pro	Ile	Ile	Ala	Glu	Tyr	Leu	Gly	Gly	Ala	Ile	Thr	Cys	260	265	270	
Asp	Ala	His	His	Ile	Thr	Asp	Pro	Arg	Ala	Asp	Gly	Leu	Gly	Val	Ser	275	280	285	
Ser	Cys	Ile	Val	Met	Ser	Leu	Val	Asp	Ala	Gly	Val	Ser	Pro	Glu	Glu	290	295	300	
Val	Asn	Tyr	Ile	Asn	Ala	His	Ala	Thr	Ser	Thr	Leu	Ala	Gly	Asp	Leu	305	310	315	320
Ala	Glu	Val	Asn	Ala	Ile	Lys	Lys	Val	Phe	Lys	Asp	Thr	Ser	Glu	Ile	325	330	335	
Lys	Met	Asn	Gly	Thr	Lys	Ser	Met	Ile	Gly	His	Gly	Leu	Gly	Ala	Ser	340	345	350	
Gly	Gly	Leu	Glu	Ala	Ile	Ala	Thr	Ile	Lys	Ala	Ile	Thr	Thr	Gly	Trp	355	360	365	
Leu	His	Pro	Thr	Ile	Asn	Gln	Tyr	Asp	Leu	Glu	Pro	Gln	Val	Thr	Ile	370	375	380	
Asp	Thr	Val	Pro	Asn	Val	Lys	Lys	Gln	His	Glu	Val	Asn	Val	Gly	Ile	385	390	395	400
Ser	Asn	Ser	Phe	Gly	Phe	Gly	Gly	His	Asn	Ser	Val	Val	Val	Phe	Ala	405	410	415	
Pro	Tyr	Lys	Pro													420			

<210> 28
 <211> 420
 <212> PRT

<213> Cuphea hookeriana

<400> 28

Lys	Lys	Lys	Pro	Ser	Ile	Lys	Gln	Arg	Arg	Val	Val	Val	Thr	Gly	Met	
1				5					10					15		
Gly	Val	Val	Thr	Pro	Leu	Gly	His	Asp	Pro	Asp	Val	Phe	Tyr	Asn	Asn	
			20					25					30			
Leu	Leu	Asp	Gly	Thr	Ser	Gly	Ile	Ser	Glu	Ile	Glu	Thr	Phe	Asp	Cys	
		35					40					45				
Ala	Gln	Phe	Pro	Thr	Arg	Ile	Ala	Gly	Glu	Ile	Lys	Ser	Phe	Ser	Thr	
	50					55					60					
Asp	Gly	Trp	Val	Ala	Pro	Lys	Leu	Ser	Lys	Arg	Met	Asp	Lys	Phe	Met	
65					70					75					80	
Leu	Tyr	Met	Leu	Thr	Ala	Gly	Lys	Lys	Ala	Leu	Thr	Asn	Gly	Gly	Ile	
				85					90					95		
Thr	Glu	Asp	Val	Met	Lys	Glu	Leu	Asp	Lys	Arg	Lys	Cys	Gly	Val	Leu	
			100					105					110			
Ile	Gly	Ser	Ala	Met	Gly	Gly	Met	Lys	Val	Phe	Asn	Asp	Ala	Ile	Glu	
		115					120					125				
Ala	Leu	Arg	Ile	Ser	Tyr	Lys	Lys	Met	Asn	Pro	Phe	Cys	Val	Pro	Phe	
	130					135					140					
Ala	Thr	Thr	Asn	Met	Gly	Ser	Ala	Met	Leu	Ala	Met	Asp	Leu	Gly	Trp	
145					150				155						160	
Met	Gly	Pro	Asn	Tyr	Ser	Ile	Ser	Thr	Ala	Cys	Ala	Thr	Ser	Asn	Phe	
				165					170					175		
Cys	Ile	Leu	Asn	Ala	Ala	Asn	His	Ile	Ile	Arg	Gly	Glu	Ala	Asp	Val	
			180					185					190			
Met	Leu	Cys	Gly	Gly	Ser	Asp	Ala	Val	Ile	Ile	Pro	Ile	Gly	Met	Gly	
		195					200					205				
Gly	Phe	Val	Ala	Cys	Arg	Ala	Leu	Ser	Gln	Arg	Asn	Ala	Asp	Pro	Thr	
	210					215					220					
Lys	Ala	Ser	Arg	Pro	Trp	Asp	Ser	Asn	Arg	Asp	Gly	Phe	Val	Met	Gly	
225					230					235					240	
Glu	Gly	Ala	Gly	Val	Leu	Leu	Leu	Glu	Glu	Leu	Glu	His	Ala	Lys	Lys	
				245					250					255		
Arg	Gly	Ala	Thr	Ile	Tyr	Ala	Glu	Phe	Leu	Gly	Gly	Ser	Phe	Thr	Cys	
			260					265					270			
Asp	Ala	Tyr	His	Met	Thr	Glu	Pro	His	Pro	Asp	Gly	Ala	Gly	Val	Ile	
		275					280					285				

Leu Cys Ile Glu Lys Ala Leu Ala Gln Ser Gly Val Ser Arg Glu Asp
 290 295 300

Val Asn Tyr Ile Asn Ala His Ala Thr Ser Thr Pro Ala Gly Asp Ile
 305 310 315 320

Lys Glu Tyr Gln Ala Leu Ile His Cys Phe Gly Gln Asn Asn Glu Leu
 325 330 335

Lys Val Asn Ser Thr Lys Ser Met Ile Gly His Leu Leu Gly Ala Ala
 340 345 350

Gly Gly Val Glu Ala Val Ser Val Val Gln Ala Ile Arg Thr Gly Trp
 355 360 365

Ile His Pro Asn Ile Asn Leu Glu Asn Pro Asp Glu Gly Val Asp Thr
 370 375 380

Lys Leu Leu Val Gly Pro Lys Lys Glu Arg Leu Asn Ile Lys Val Gly
 385 390 395 400

Leu Ser Asn Ser Phe Gly Phe Gly Gly His Asn Ser Ser Ile Leu Phe
 405 410 415

Ala Pro Tyr Asn
 420

<210> 29
 <211> 420
 <212> PRT
 <213> Cuphea hookeriana

<400> 29

Asn Lys Lys Pro Ala Thr Lys Gln Arg Arg Val Val Val Thr Gly Met
 1 5 10 15

Gly Val Val Thr Pro Leu Gly His Asp Pro Asp Val Tyr Tyr Asn Asn
 20 25 30

Leu Leu Asp Gly Ile Ser Gly Ile Ser Glu Ile Glu Asn Phe Asp Cys
 35 40 45

Ser Gln Phe Pro Thr Arg Ile Ala Gly Glu Ile Lys Ser Phe Ser Thr
 50 55 60

Asp Gly Trp Val Ala Pro Lys Phe Ser Glu Arg Met Asp Lys Phe Met
 65 70 75 80

Leu Tyr Met Leu Thr Ala Gly Lys Lys Ala Leu Ala Asp Gly Gly Ile
 85 90 95

Thr Glu Asp Ala Met Lys Glu Leu Asn Lys Arg Lys Cys Gly Val Leu
 100 105 110

Ile Gly Ser Gly Leu Gly Gly Met Lys Val Phe Ser Asp Ser Ile Glu

115					120					125					
Ala	Leu	Arg	Thr	Ser	Tyr	Lys	Lys	Ile	Ser	Pro	Phe	Cys	Val	Pro	Phe
130						135					140				
Ser	Thr	Thr	Asn	Met	Gly	Ser	Ala	Ile	Leu	Ala	Met	Asp	Leu	Gly	Trp
145					150					155					160
Met	Gly	Pro	Asn	Tyr	Ser	Ile	Ser	Thr	Ala	Cys	Ala	Thr	Ser	Asn	Phe
				165					170					175	
Cys	Ile	Leu	Asn	Ala	Ala	Asn	His	Ile	Ile	Lys	Gly	Glu	Ala	Asp	Met
			180					185						190	
Met	Leu	Cys	Gly	Gly	Ser	Asp	Ala	Ala	Val	Leu	Pro	Val	Gly	Leu	Gly
	195						200					205			
Gly	Phe	Val	Ala	Cys	Arg	Ala	Leu	Ser	Gln	Arg	Asn	Asn	Asp	Pro	Thr
210						215					220				
Lys	Ala	Ser	Arg	Pro	Trp	Asp	Ser	Asn	Arg	Asp	Gly	Phe	Val	Met	Gly
225					230					235					240
Glu	Gly	Ala	Gly	Val	Leu	Leu	Leu	Glu	Glu	Leu	Glu	His	Ala	Lys	Lys
				245					250					255	
Arg	Gly	Ala	Thr	Ile	Tyr	Ala	Glu	Phe	Leu	Gly	Gly	Ser	Phe	Thr	Cys
			260					265						270	
Asp	Ala	Tyr	His	Met	Thr	Glu	Pro	His	Pro	Glu	Gly	Ala	Gly	Val	Ile
		275					280					285			
Leu	Cys	Ile	Glu	Lys	Ala	Leu	Ala	Gln	Ser	Gly	Val	Ser	Arg	Glu	Asp
	290					295					300				
Val	Asn	Tyr	Ile	Asn	Ala	His	Ala	Thr	Ser	Thr	Pro	Ala	Gly	Asp	Ile
305					310					315					320
Lys	Glu	Tyr	Gln	Ala	Leu	Ala	His	Cys	Phe	Gly	Gln	Asn	Ser	Glu	Leu
				325					330					335	
Arg	Val	Asn	Ser	Thr	Lys	Ser	Met	Ile	Gly	His	Leu	Leu	Gly	Gly	Ala
			340					345					350		
Gly	Gly	Val	Glu	Ala	Val	Ala	Val	Val	Gln	Ala	Ile	Arg	Thr	Gly	Trp
		355					360					365			
Ile	His	Pro	Asn	Ile	Asn	Leu	Glu	Asp	Pro	Asp	Glu	Gly	Val	Asp	Ala
	370					375					380				
Lys	Leu	Leu	Val	Gly	Pro	Lys	Lys	Glu	Lys	Leu	Lys	Val	Lys	Val	Gly
385					390					395					400
Leu	Ser	Asn	Ser	Phe	Gly	Phe	Gly	Gly	His	Asn	Ser	Ser	Ile	Leu	Phe
				405					410					415	
Ala	Pro	Cys	Asn												

420

<210> 30
 <211> 420
 <212> PRT
 <213> Cuphea pullcherima

<400> 30

Lys	Lys	Lys	Pro	Ser	Ile	Lys	Gln	Arg	Arg	Val	Val	Val	Thr	Gly	Met
1				5					10					15	
Gly	Val	Val	Thr	Pro	Leu	Gly	His	Asp	Pro	Asp	Val	Phe	Tyr	Asn	Asn
			20					25					30		
Leu	Leu	Asp	Gly	Thr	Ser	Gly	Ile	Ser	Glu	Ile	Glu	Thr	Phe	Asp	Cys
		35					40					45			
Ala	Gln	Phe	Pro	Thr	Arg	Ile	Ala	Gly	Glu	Ile	Lys	Ser	Phe	Ser	Thr
	50					55					60				
Asp	Gly	Trp	Val	Ala	Pro	Lys	Leu	Ser	Lys	Arg	Met	Asp	Lys	Phe	Met
65					70					75					80
Leu	Tyr	Met	Leu	Thr	Ala	Gly	Lys	Lys	Ala	Leu	Thr	Asp	Gly	Gly	Ile
				85					90					95	
Thr	Glu	Asp	Val	Met	Lys	Glu	Leu	Asp	Lys	Arg	Lys	Cys	Gly	Val	Leu
			100					105					110		
Ile	Gly	Ser	Ala	Met	Gly	Gly	Met	Lys	Val	Phe	Asn	Asp	Ala	Ile	Glu
		115					120					125			
Ala	Leu	Arg	Ile	Ser	Tyr	Lys	Lys	Met	Asn	Pro	Phe	Cys	Val	Pro	Phe
	130					135					140				
Ala	Thr	Thr	Asn	Met	Gly	Ser	Ala	Met	Leu	Ala	Met	Asp	Leu	Gly	Trp
145					150					155					160
Met	Gly	Pro	Asn	Tyr	Ser	Ile	Ser	Thr	Ala	Cys	Ala	Thr	Ser	Asn	Phe
				165					170					175	
Cys	Ile	Met	Asn	Ala	Ala	Asn	His	Ile	Ile	Arg	Gly	Glu	Ala	Asp	Val
			180					185					190		
Met	Leu	Cys	Gly	Gly	Ser	Asp	Ala	Val	Ile	Ile	Pro	Ile	Gly	Met	Gly
	195						200					205			
Gly	Phe	Val	Ala	Cys	Arg	Ala	Leu	Ser	Gln	Arg	Asn	Ser	Asp	Pro	Thr
	210					215					220				
Lys	Ala	Ser	Arg	Pro	Trp	Asp	Ser	Asn	Arg	Asp	Gly	Phe	Val	Met	Gly
225					230					235					240
Glu	Gly	Ala	Gly	Val	Leu	Leu	Leu	Glu	Glu	Leu	Glu	His	Ala	Lys	Lys
				245				250						255	

Arg Gly Ala Thr Ile Tyr Ala Glu Phe Leu Gly Gly Ser Phe Thr Cys
260 265 270

Asp Ala Tyr His Met Thr Glu Pro His Pro Asp Gly Ala Gly Val Ile
275 280 285

Leu Cys Ile Glu Lys Ala Leu Ala Gln Ser Gly Val Ser Arg Glu Asp
290 295 300

Val Asn Tyr Ile Asn Ala His Ala Thr Ser Thr Pro Ala Gly Asp Ile
305 310 315 320

Lys Glu Tyr Gln Ala Leu Ile His Cys Phe Gly Gln Asn Arg Glu Leu
325 330 335

Lys Val Asn Ser Thr Lys Ser Met Ile Gly His Leu Leu Gly Ala Ala
340 345 350

Gly Gly Val Glu Ala Val Ser Val Val Gln Ala Ile Arg Thr Gly Trp
355 360 365

Ile His Pro Asn Ile Asn Leu Glu Asn Pro Asp Glu Gly Val Asp Thr
370 375 380

Lys Leu Leu Val Gly Pro Lys Lys Glu Arg Leu Asn Val Lys Val Gly
385 390 395 400

Leu Ser Asn Ser Phe Gly Phe Gly Gly His Asn Ser Ser Ile Leu Phe
405 410 415

Ala Pro Tyr Ile
420

<210> 31
<211> 421
<212> PRT
<213> Cuphea wrightii

<400> 31

Lys Lys Lys Pro Val Ile Lys Gln Arg Arg Val Val Val Thr Gly Met
1 5 10 15

Gly Val Val Thr Pro Leu Gly His Glu Pro Asp Val Phe Tyr Asn Asn
20 25 30

Leu Leu Asp Gly Val Ser Gly Ile Ser Glu Ile Glu Thr Phe Asp Cys
35 40 45

Thr Gln Phe Pro Thr Arg Ile Ala Gly Glu Ile Lys Ser Phe Ser Thr
50 55 60

Asp Gly Trp Val Ala Pro Lys Leu Ser Lys Arg Met Asp Lys Phe Met
65 70 75 80

Leu Tyr Leu Leu Thr Ala Gly Lys Lys Ala Leu Ala Asp Gly Gly Ile
85 90 95

Thr	Asp	Glu	Val	Met	Lys	Glu	Leu	Asp	Lys	Arg	Lys	Cys	Gly	Val	Leu	100	105	110
Ile	Gly	Ser	Gly	Met	Gly	Gly	Met	Lys	Val	Phe	Asn	Asp	Ala	Ile	Glu	115	120	125
Ala	Leu	Arg	Val	Ser	Tyr	Lys	Lys	Met	Asn	Pro	Phe	Cys	Val	Pro	Phe	130	135	140
Ala	Thr	Thr	Asn	Met	Gly	Ser	Ala	Met	Leu	Ala	Met	Asp	Leu	Gly	Trp	145	150	155
Met	Gly	Pro	Asn	Tyr	Ser	Ile	Ser	Thr	Ala	Cys	Ala	Thr	Ser	Asn	Phe	165	170	175
Cys	Ile	Leu	Asn	Ala	Ala	Asn	His	Ile	Ile	Arg	Gly	Glu	Ala	Asp	Met	180	185	190
Met	Leu	Cys	Gly	Gly	Ser	Asp	Ala	Val	Ile	Ile	Pro	Ile	Gly	Leu	Gly	195	200	205
Gly	Phe	Val	Ala	Cys	Arg	Ala	Leu	Ser	Gln	Arg	Asn	Ser	Asp	Pro	Thr	210	215	220
Lys	Ala	Ser	Arg	Pro	Trp	Asp	Ser	Asn	Arg	Asp	Gly	Phe	Val	Met	Gly	225	230	235
Glu	Gly	Ala	Gly	Val	Leu	Leu	Leu	Glu	Glu	Leu	Glu	His	Ala	Lys	Lys	245	250	255
Arg	Gly	Ala	Thr	Ile	Tyr	Ala	Glu	Phe	Leu	Gly	Gly	Ser	Phe	Thr	Cys	260	265	270
Asp	Ala	Tyr	His	Met	Thr	Glu	Pro	His	Pro	Glu	Gly	Ala	Gly	Val	Ile	275	280	285
Leu	Cys	Ile	Glu	Lys	Ala	Leu	Ala	Gln	Ala	Gly	Val	Ser	Lys	Glu	Asp	290	295	300
Val	Asn	Tyr	Ile	Asn	Ala	His	Ala	Thr	Ser	Thr	Ser	Ala	Gly	Asp	Ile	305	310	315
Lys	Glu	Tyr	Gln	Ala	Leu	Ala	Arg	Cys	Phe	Gly	Gln	Asn	Ser	Glu	Leu	325	330	335
Arg	Val	Asn	Ser	Thr	Lys	Ser	Met	Ile	Gly	His	Leu	Leu	Gly	Ala	Ala	340	345	350
Gly	Gly	Val	Glu	Ala	Val	Thr	Val	Val	Gln	Ala	Ile	Arg	Thr	Gly	Trp	355	360	365
Ile	His	Pro	Asn	Leu	Asn	Leu	Glu	Asp	Pro	Asp	Lys	Ala	Val	Asp	Ala	370	375	380
Lys	Leu	Leu	Val	Gly	Pro	Lys	Lys	Glu	Arg	Leu	Asn	Val	Lys	Val	Gly	385	390	395
																		400

Leu Ser Asn Ser Phe Gly Phe Gly Gly His Asn Ser Ser Ile Leu Phe
 405 410 415

Ala Pro Cys Asn Val
 420

<210> 32
 <211> 420
 <212> PRT
 <213> Cuphea wrightii

<400> 32

Lys Lys Lys Pro Val Thr Lys Gln Arg Arg Val Val Val Thr Gly Met
 1 5 10 15

Gly Val Val Thr Pro Leu Gly His Asp Pro Asp Val Phe Tyr Asn Asn
 20 25 30

Leu Leu Asp Gly Val Ser Gly Ile Ser Glu Ile Glu Thr Phe Asp Cys
 35 40 45

Thr Gln Phe Pro Thr Arg Ile Ala Gly Glu Ile Lys Ser Phe Ser Thr
 50 55 60

Asp Gly Trp Val Ala Pro Lys Leu Ser Lys Arg Met Asp Lys Phe Met
 65 70 75 80

Leu Tyr Met Leu Thr Ala Gly Lys Lys Ala Leu Ala Asp Ala Gly Ile
 85 90 95

Thr Glu Asp Val Met Lys Glu Leu Asp Lys Arg Lys Cys Gly Val Leu
 100 105 110

Ile Gly Ser Gly Met Gly Gly Met Lys Leu Phe Asn Asp Ser Ile Glu
 115 120 125

Ala Leu Arg Ile Ser Tyr Lys Lys Met Asn Pro Phe Cys Val Pro Phe
 130 135 140

Ala Thr Thr Asn Met Gly Ser Ala Met Leu Ala Met Asp Leu Gly Trp
 145 150 155 160

Met Gly Pro Asn Tyr Ser Ile Ser Thr Ala Cys Ala Thr Ser Asn Phe
 165 170 175

Cys Ile Leu Asn Ala Ala Asn His Ile Ile Arg Gly Glu Ala Asp Met
 180 185 190

Met Leu Cys Gly Gly Ser Asp Ala Ala Ile Ile Pro Ile Gly Leu Gly
 195 200 205

Gly Phe Val Ala Cys Arg Ala Leu Ser Gln Arg Asn Asn Asp Pro Thr
 210 215 220

Lys Ala Ser Arg Pro Trp Asp Ser Asn Arg Asp Gly Phe Val Met Gly

225		230		235		240
Glu Gly Ala Gly Val Leu Leu Leu Glu Glu Leu Glu His Ala Lys Lys						
	245		250		255	
Arg Gly Ala Thr Ile Tyr Ala Glu Phe Leu Gly Gly Ser Phe Thr Cys						
	260		265		270	
Asp Ala Tyr His Met Thr Glu Pro His Pro Glu Gly Ala Gly Val Ile						
	275		280		285	
Leu Cys Ile Glu Arg Ala Leu Ala Gln Ser Gly Val Ser Lys Glu Asp						
	290		295		300	
Val Asn Tyr Ile Asn Ala His Ala Thr Ser Thr Pro Ala Gly Asp Ile						
305		310		315		320
Lys Glu Tyr Gln Ala Leu Ala Arg Ile Phe Ser Gln Asn Ser Glu Leu						
	325		330		335	
Arg Val Asn Ser Thr Lys Ser Met Ile Gly His Leu Leu Gly Ala Ala						
	340		345		350	
Gly Gly Val Glu Ala Val Thr Val Val Gln Ala Ile Arg Thr Gly Trp						
	355		360		365	
Ile His Pro Asn Ile Asn Leu Glu Asn Pro Asp Asp Gly Val Asp Ala						
	370		375		380	
Lys Leu Leu Val Gly Pro Lys Lys Glu Lys Leu Lys Val Lys Val Gly						
385		390		395		400
Leu Ser Asn Ser Phe Gly Phe Gly Gly His Asn Ser Ser Ile Leu Phe						
	405		410		415	
Ala Pro Cys Asn						
	420					

<210> 33
 <211> 420
 <212> PRT
 <213> Hordeum vulgare

<400> 33

Lys Lys Arg Pro Asp Val Lys Gln Arg Arg Val Val Val Thr Gly Met						
1		5		10		15
Gly Val Val Thr Pro Leu Gly His Asp Pro Asp Val Phe Tyr Thr Asn						
	20		25		30	
Leu Leu Asp Gly His Ser Gly Ile Ser Glu Ile Glu Thr Phe Asp Cys						
	35		40		45	
Ser Lys Phe Pro Thr Arg Ile Ala Gly Glu Ile Lys Ser Phe Ser Thr						
	50		55		60	

Glu	Gly	Trp	Val	Val	Pro	Lys	Leu	Ser	Lys	Arg	Met	Asp	Lys	Phe	Met	65	70	75	80
Leu	Tyr	Leu	Ile	Thr	Ala	Gly	Lys	Lys	Ala	Leu	Glu	Asn	Gly	Gly	Leu	85	90	95	
Thr	Glu	Glu	Val	Arg	Asn	Glu	Leu	Asp	Lys	Thr	Arg	Cys	Gly	Val	Leu	100	105	110	
Ile	Gly	Ser	Ala	Met	Gly	Gly	Met	Lys	Val	Phe	Asn	Asp	Ala	Ile	Glu	115	120	125	
Ala	Leu	Arg	Val	Ser	Tyr	Arg	Lys	Met	Asn	Pro	Phe	Cys	Val	Pro	Phe	130	135	140	
Ala	Thr	Thr	Asn	Met	Gly	Ser	Ala	Ile	Leu	Ala	Met	Asp	Leu	Gly	Trp	145	150	155	160
Met	Gly	Pro	Asn	Tyr	Ser	Ile	Ser	Thr	Ala	Cys	Ala	Thr	Ser	Asn	Phe	165	170	175	
Cys	Ile	Leu	Asn	Ala	Ala	Asn	His	Ile	Arg	Arg	Gly	Glu	Ala	Asp	Val	180	185	190	
Met	Leu	Cys	Gly	Gly	Ser	Asp	Ala	Pro	Leu	Ile	Pro	Ile	Gly	Leu	Gly	195	200	205	
Gly	Phe	Val	Ala	Cys	Arg	Ala	Leu	Ser	Gln	Arg	Asn	Ser	Asp	Pro	Thr	210	215	220	
Lys	Ala	Ser	Arg	Pro	Trp	Asp	Met	Asp	Arg	Asp	Gly	Phe	Val	Met	Gly	225	230	235	240
Glu	Gly	Ala	Gly	Val	Leu	Val	Leu	Glu	Glu	Leu	Glu	His	Ala	Lys	Gln	245	250	255	
Arg	Gly	Ala	Thr	Ile	Tyr	Ala	Glu	Phe	Leu	Gly	Gly	Ser	Phe	Thr	Cys	260	265	270	
Asp	Ala	Tyr	His	Met	Thr	Glu	Pro	His	Pro	Glu	Gly	Thr	Gly	Ile	Thr	275	280	285	
Leu	Cys	Ile	Glu	Lys	Ala	Leu	Ala	Asp	Ser	Gly	Val	Ala	Arg	Glu	Glu	290	295	300	
Ile	Asn	Tyr	Val	Asn	Ala	His	Ala	Thr	Ser	Thr	Gln	Ser	Gly	Asp	Leu	305	310	315	320
Lys	Glu	Tyr	Glu	Ala	Ile	Val	Arg	Cys	Phe	Gly	Gln	Asn	Pro	Gln	Leu	325	330	335	
Arg	Val	Asn	Ser	Thr	Lys	Ser	Met	Thr	Gly	His	Leu	Ile	Gly	Ala	Ala	340	345	350	
Gly	Gly	Ile	Glu	Ala	Val	Ala	Cys	Val	Gln	Ala	Ile	Arg	Thr	Gly	Trp	355	360	365	

Val His Pro Asn Leu Asn Leu Glu Asn Pro Glu Lys Val Val Asp Val
 370 375 380

Gly Val Leu Val Gly Ser Glu Lys Glu Arg Cys Glu Val Lys Val Ala
 385 390 395 400

Leu Ser Asn Ser Phe Gly Phe Gly Gly His Asn Ser Ser Ile Leu Phe
 405 410 415

Ala Pro Phe Lys
 420

<210> 34

<211> 419

<212> PRT

<213> Hordeum vulgare

<400> 34

Asn Asn Lys Ser Glu Thr Lys Gln Arg Arg Val Val Val Thr Gly Met
 1 5 10 15

Gly Val Val Thr Pro Leu Gly His Glu Pro Asp Glu Phe Tyr Asn Asn
 20 25 30

Leu Leu Gln Gly Val Ser Gly Val Ser Glu Ile Glu Ala Phe Asp Cys
 35 40 45

Ser Ser Tyr Pro Thr Arg Ile Ala Gly Glu Ile Lys Ser Phe Ser Thr
 50 55 60

Asp Gly Trp Val Ala Pro Lys Leu Ala Lys Arg Met Asp Lys Phe Met
 65 70 75 80

Gln Tyr Leu Ile Val Ala Gly Lys Lys Ala Leu Asp Asn Gly Gly Val
 85 90 95

Thr Glu Asp Ile Met Asn Glu Leu Asp Lys Ser Arg Cys Gly Val Leu
 100 105 110

Ile Gly Ser Gly Met Gly Gly Met Lys Val Phe Ser Asp Ala Ile Glu
 115 120 125

Ala Leu Arg Val Ser Tyr Arg Lys Met Asn Pro Phe Cys Val Pro Phe
 130 135 140

Ala Thr Thr Asn Met Gly Ser Ala Val Leu Ala Met Asp Leu Gly Trp
 145 150 155 160

Met Gly Pro Asn Tyr Ser Ile Ser Thr Ala Cys Ala Thr Ser Asn Phe
 165 170 175

Cys Ile Leu Ser Ala Ala Asn His Ile Met Arg Gly Glu Thr Asp Leu
 180 185 190

Met Leu Cys Gly Gly Ser Asp Ala Pro Ile Ile Pro Ile Gly Leu Gly
 195 200 205

Gly Phe Val Ala Cys Arg Ala Leu Ser Gln Arg Asn Ser Asp Pro Thr
 210 215 220
 Lys Ala Ser Arg Pro Trp Asp Val Asp Arg Asp Gly Phe Val Met Gly
 225 230 235 240
 Glu Gly Ala Gly Val Leu Leu Leu Glu Glu Leu Glu His Ala Lys Gln
 245 250 255
 Arg Gly Ala Glu Ile Tyr Ala Glu Phe Leu Gly Gly Asn Phe Thr Cys
 260 265 270
 Asp Ala Tyr His Met Thr Glu Pro His Pro Glu Gly Lys Gly Val Ile
 275 280 285
 Leu Cys Val Glu Asn Ala Leu Ala Asp Ala Gly Val Thr Arg Gln Asp
 290 295 300
 Ile Asn Tyr Val Asn Ala His Ala Thr Ser Thr Gln Leu Gly Asp Leu
 305 310 315 320
 Lys Glu Phe Glu Ala Leu Arg Arg Cys Phe Gly Gln Asn Pro Gln Leu
 325 330 335
 Arg Val Asn Ser Thr Lys Ser Met Thr Gly His Leu Leu Gly Ala Ala
 340 345 350
 Gly Gly Ile Glu Ala Val Ala Ala Ile Gln Ala Ile Arg Thr Gly Trp
 355 360 365
 Ile His Pro Asn Ile Asn Leu Asn Asn Pro Glu Lys Asn Val Asp Val
 370 375 380
 Ser Leu Leu Val Gly Ser Gln Lys Glu Arg Cys Asp Val Lys Val Ala
 385 390 395 400
 Leu Ser Asn Ser Phe Gly Phe Gly Gly His Asn Ser Ser Ile Leu Phe
 405 410 415

Ala Pro Phe

<210> 35
 <211> 420
 <212> PRT
 <213> Ricinus communis

<400> 35

Asn Lys Lys Pro Leu Met Lys Gln Arg Arg Val Val Val Thr Gly Met
 1 5 10 15
 Gly Val Val Ser Pro Leu Gly His Asp Ile Asp Val Tyr Tyr Asn Asn
 20 25 30
 Leu Leu Asp Gly Ser Ser Gly Ile Ser Gln Ile Asp Ser Phe Asp Cys

35					40					45					
Ala	Gln	Phe	Pro	Thr	Arg	Ile	Ala	Gly	Glu	Ile	Lys	Ser	Phe	Ser	Thr
50						55					60				
Asp	Gly	Trp	Val	Ala	Pro	Lys	Leu	Ser	Lys	Arg	Met	Asp	Lys	Phe	Met
65					70					75					80
Leu	Tyr	Met	Leu	Thr	Ala	Gly	Lys	Lys	Ala	Leu	Ala	Asp	Gly	Gly	Ile
				85					90					95	
Thr	Glu	Asp	Met	Met	Asp	Glu	Leu	Asp	Lys	Ala	Arg	Cys	Gly	Val	Leu
			100					105					110		
Ile	Gly	Ser	Ala	Met	Gly	Gly	Met	Lys	Val	Phe	Asn	Asp	Ala	Ile	Glu
		115					120					125			
Ala	Leu	Arg	Ile	Ser	Tyr	Arg	Lys	Met	Asn	Pro	Phe	Cys	Val	Pro	Phe
	130					135					140				
Ala	Thr	Thr	Asn	Met	Gly	Ser	Ala	Met	Leu	Ala	Met	Asp	Leu	Gly	Trp
145					150					155					160
Met	Gly	Pro	Asn	Tyr	Ser	Ile	Ser	Thr	Ala	Cys	Ala	Thr	Ser	Asn	Phe
				165					170					175	
Cys	Ile	Leu	Asn	Ala	Ala	Asn	His	Ile	Ile	Arg	Gly	Glu	Ala	Asp	Ile
			180					185					190		
Met	Leu	Cys	Gly	Gly	Ser	Asp	Ala	Ala	Ile	Ile	Pro	Ile	Gly	Leu	Gly
		195					200					205			
Gly	Phe	Val	Ala	Cys	Arg	Ala	Leu	Ser	Gln	Arg	Asn	Asp	Asp	Pro	Thr
	210					215					220				
Lys	Ala	Ser	Arg	Pro	Trp	Asp	Met	Asn	Arg	Asp	Gly	Phe	Val	Met	Gly
225					230					235					240
Glu	Gly	Ala	Gly	Val	Leu	Leu	Leu	Glu	Glu	Leu	Glu	His	Ala	Lys	Lys
				245				250						255	
Arg	Gly	Ala	Asn	Ile	Tyr	Ala	Glu	Phe	Leu	Gly	Gly	Ser	Phe	Thr	Cys
			260					265					270		
Asp	Ala	Tyr	His	Met	Thr	Glu	Pro	Arg	Pro	Asp	Gly	Val	Gly	Val	Ile
		275					280					285			
Leu	Cys	Ile	Glu	Lys	Ala	Leu	Ala	Arg	Ser	Gly	Val	Ser	Lys	Glu	Glu
	290					295					300				
Val	Asn	Tyr	Ile	Asn	Ala	His	Ala	Thr	Ser	Thr	Pro	Ala	Gly	Asp	Leu
305					310					315					320
Lys	Glu	Tyr	Glu	Ala	Leu	Met	Arg	Cys	Phe	Ser	Gln	Asn	Pro	Asp	Leu
				325					330					335	
Arg	Val	Asn	Ser	Thr	Lys	Ser	Met	Ile	Gly	His	Leu	Leu	Gly	Ala	Ala

340 345 350
 Gly Ala Val Glu Ala Ile Ala Thr Ile Gln Ala Ile Arg Thr Gly Trp
 355 360 365
 Val His Pro Asn Ile Asn Leu Glu Asn Pro Glu Glu Gly Val Asp Thr
 370 375 380
 Lys Val Leu Val Gly Pro Lys Lys Glu Arg Leu Asp Ile Lys Val Ala
 385 390 395 400
 Leu Ser Asn Ser Phe Gly Phe Gly Gly His Asn Ser Ser Ile Ile Phe
 405 410 415
 Ala Pro Tyr Lys
 420

 <210> 36
 <211> 413
 <212> PRT
 <213> Caenorhabditis elegans

 <220>
 <221> misc_feature
 <222> (53)..(53)
 <223> Xaa in position 53 in unknown.

 <400> 36

 Met Lys Leu Lys Ile Asn Lys Asn Phe Glu Met His Arg Val Val Ile
 1 5 10 15
 Thr Gly Met Gly Ala Ile Ser Pro Phe Gly Val Thr Val Asn Ala Leu
 20 25 30
 Arg Asn Gly Leu Asn Glu Gly Arg Ser Gly Leu Lys Tyr Asp Glu Ile
 35 40 45
 Leu Lys Phe Val Xaa Gly Ala Val Pro Gly Glu Arg Val Glu Asp Arg
 50 55 60
 Trp Ser Thr Gly Gln Gln Arg Glu Met Ser Lys Ala Ser Met Phe Val
 65 70 75 80
 Leu Ala Ala Ser Glu Glu Ala Leu Lys Gln Ala Lys Ala Glu Asp Val
 85 90 95
 Asp His Asn Glu Thr Leu Val Asn Ile Gly Thr Cys Met Ser Asp Leu
 100 105 110
 Glu His Ile Gly Glu Thr Ala Gln Lys Val Ser Glu Gly Gln Ser Arg
 115 120 125
 Arg Val Ser Pro Tyr Phe Val Pro Arg Ile Leu Asn Asn Leu Pro Ala
 130 135 140

Gly	Tyr	Val	Ala	Met	Lys	Tyr	Lys	Met	Arg	Gly	Gly	Val	Glu	Ser	Thr	145	150	155	160
Ser	Thr	Ala	Cys	Ala	Thr	Gly	Leu	His	Cys	Ile	Gly	Asn	Ser	Phe	Arg	165	170	175	
Ser	Ile	Arg	Tyr	Gly	Asp	Ser	Arg	Arg	Ala	Leu	Ala	Gly	Ala	Val	Glu	180	185	190	
Cys	Ala	Leu	Asn	Pro	Ile	Ala	Leu	Ala	Gly	Phe	Asp	Arg	Met	Arg	Ala	195	200	205	
Leu	Ala	Arg	Gly	Asp	Gln	Pro	Asn	Ile	Ser	Arg	Pro	Phe	Asp	Lys	Lys	210	215	220	
Arg	Ala	Gly	Phe	Val	Met	Ser	Glu	Gly	Val	Gly	Leu	Val	Phe	Met	Glu	225	230	235	240
Arg	Leu	Glu	Asp	Ala	Gln	Ala	Arg	Gly	Ala	Gln	Ile	Leu	Ala	Glu	Val	245	250	255	
Val	Gly	Tyr	Gly	Ile	Ser	Ser	Asp	Cys	Tyr	His	Ile	Ser	Thr	Pro	Asp	260	265	270	
Pro	Ser	Ala	Ile	Gly	Ala	Val	Leu	Ser	Met	Asn	Arg	Ala	Ile	Gly	Asn	275	280	285	
Ala	His	Leu	Glu	Pro	Lys	Asp	Ile	Gly	Tyr	Val	Asn	Ala	His	Ala	Thr	290	295	300	
Ser	Thr	Pro	Asn	Gly	Asp	Ser	Val	Glu	Ala	Glu	Ala	Val	Arg	Gln	Val	305	310	315	320
Phe	Pro	Glu	Gln	Asn	Ile	Ala	Val	Ser	Ser	Val	Lys	Gly	His	Ile	Gly	325	330	335	
His	Leu	Leu	Gly	Ala	Ala	Gly	Ser	Val	Glu	Ala	Ile	Ala	Thr	Ile	Phe	340	345	350	
Ala	Met	Asn	Asp	Asp	Val	Leu	Pro	Ala	Asn	Arg	Asn	Leu	Glu	Glu	Thr	355	360	365	
Asp	Glu	Gly	Asn	Gly	Leu	Asn	Leu	Leu	Arg	Glu	Asn	Gln	Lys	Trp	Ser	370	375	380	
Asp	Val	Ser	Gly	Asn	Lys	Ser	Arg	Ile	Ser	Ile	Cys	Asn	Ser	Phe	Gly	385	390	395	400
Phe	Gly	Ala	Thr	Asn	Ala	Ser	Leu	Ile	Leu	Lys	Gln	Phe	405	410					

<210> 37
 <211> 442
 <212> PRT
 <213> *Saccharomyces cerevisiae*
 <400> 37

Met	Ser	Arg	Arg	Val	Val	Ile	Thr	Gly	Leu	Gly	Cys	Val	Thr	Pro	Leu	
1				5					10					15		
Gly	Arg	Ser	Leu	Ser	Glu	Ser	Trp	Gly	Asn	Leu	Leu	Ser	Ser	Lys	Asn	
			20					25					30			
Gly	Leu	Thr	Pro	Ile	Thr	Ser	Leu	Pro	Asn	Tyr	Asn	Glu	Asp	Tyr	Lys	
		35					40					45				
Leu	Arg	Glu	Lys	Ser	Ile	Pro	Ser	Thr	Ile	Thr	Val	Gly	Lys	Ile	Pro	
	50					55					60					
Glu	Asn	Phe	Gln	Asn	Glu	Asn	Ser	Ala	Ile	Asn	Lys	Leu	Leu	Phe	Thr	
65					70					75					80	
Ser	Gln	Asp	Glu	Arg	Arg	Thr	Ser	Ser	Phe	Ile	Lys	Leu	Ala	Leu	Arg	
				85					90					95		
Thr	Thr	Tyr	Glu	Ala	Leu	His	Asn	Ala	Gly	Leu	Leu	Asn	Pro	Asn	Asp	
			100					105					110			
Ile	Thr	Ile	Asn	Thr	Ser	Leu	Cys	Asn	Leu	Asp	His	Phe	Gly	Cys	Leu	
		115					120					125				
Ile	Gly	Ser	Gly	Ile	Gly	Ser	Ile	Gln	Asp	Ile	Tyr	Gln	Thr	Ser	Leu	
	130					135					140					
Gln	Phe	His	Asn	Asp	Asn	Lys	Arg	Ile	Asn	Pro	Tyr	Phe	Val	Pro	Lys	
145					150					155					160	
Ile	Leu	Thr	Asn	Met	Ala	Ala	Gly	Asn	Val	Ser	Ile	Lys	Phe	Asn	Leu	
				165				170						175		
Arg	Gly	Leu	Ser	His	Ser	Val	Ser	Thr	Ala	Cys	Ala	Thr	Gly	Asn	Asn	
			180					185					190			
Ser	Ile	Gly	Asp	Ala	Phe	Asn	Phe	Ile	Arg	Leu	Gly	Met	Gln	Asp	Ile	
		195					200					205				
Cys	Val	Ala	Gly	Ala	Ser	Glu	Thr	Ser	Leu	His	Pro	Leu	Ser	Leu	Ala	
	210					215					220					
Gly	Phe	Ile	Arg	Ala	Lys	Ser	Ile	Thr	Thr	Asn	Gly	Ile	Ser	Arg	Pro	
225					230					235					240	
Phe	Asp	Thr	Gln	Arg	Ser	Gly	Phe	Val	Leu	Gly	Glu	Gly	Cys	Gly	Met	
				245					250					255		
Ile	Val	Met	Glu	Ser	Leu	Glu	His	Ala	Gln	Lys	Arg	Asn	Ala	Asn	Ile	
		260						265					270			
Ile	Ser	Glu	Leu	Val	Gly	Tyr	Gly	Leu	Ser	Ser	Asp	Ala	Cys	His	Ile	
		275					280					285				
Thr	Ser	Pro	Pro	Ala	Asp	Gly	Asn	Gly	Ala	Lys	Arg	Ala	Ile	Glu	Met	
		290				295					300					

Ala Leu Lys Met Ala Arg Leu Glu Pro Thr Asp Val Asp Tyr Val Asn
 305 310 315 320

Ala His Ala Thr Ser Thr Leu Leu Gly Asp Lys Ala Glu Cys Leu Ala
 325 330 335

Val Ala Ser Ala Leu Leu Pro Gly Arg Ser Lys Ser Lys Pro Leu Tyr
 340 345 350

Ile Ser Ser Asn Lys Gly Ala Ile Gly His Leu Leu Gly Ala Ala Gly
 355 360 365

Ala Val Glu Ser Ile Phe Thr Ile Cys Ser Leu Lys Asp Asp Lys Met
 370 375 380

Pro His Thr Leu Asn Leu Asp Asn Val Leu Thr Leu Glu Asn Asn Glu
 385 390 395 400

Ala Asp Lys Leu His Phe Ile Arg Asp Lys Pro Ile Val Gly Ala Asn
 405 410 415

Pro Lys Tyr Ala Leu Cys Asn Ser Phe Gly Phe Gly Gly Val Asn Thr
 420 425 430

Ser Leu Leu Phe Lys Lys Trp Glu Gly Ser
 435 440

<210> 38

<211> 410

<212> PRT

<213> Escherichia coli

<400> 38

Met Ser Lys Arg Arg Val Val Val Thr Gly Leu Gly Met Leu Ser Pro
 1 5 10 15

Val Gly Asn Thr Val Glu Ser Thr Trp Lys Ala Leu Leu Ala Gly Gln
 20 25 30

Ser Gly Ile Ser Leu Ile Asp His Phe Asp Thr Ser Ala Tyr Ala Thr
 35 40 45

Lys Phe Ala Gly Leu Val Lys Asp Phe Asn Cys Glu Asp Ile Ile Ser
 50 55 60

Arg Lys Glu Gln Arg Lys Met Asp Ala Phe Ile Gln Tyr Gly Ile Val
 65 70 75 80

Ala Gly Val Gln Ala Met Gln Asp Ser Gly Leu Glu Ile Thr Glu Glu
 85 90 95

Asn Ala Thr Arg Ile Gly Ala Ala Ile Gly Ser Gly Ile Gly Gly Leu
 100 105 110

Gly Leu Ile Glu Glu Asn His Thr Ser Leu Met Asn Gly Gly Pro Arg

115	120	125
Lys Ile Ser Pro Phe Phe Val Pro Ser Thr Ile Val Asn Met Val Ala 130 135 140		
Gly His Leu Thr Ile Met Tyr Gly Leu Arg Gly Pro Ser Ile Ser Ile 145 150 155 160		
Ala Thr Ala Cys Thr Ser Gly Val His Asn Ile Gly His Ala Ala Arg 165 170 175		
Ile Ile Ala Tyr Gly Asp Ala Asp Val Met Val Ala Gly Gly Ala Glu 180 185 190		
Lys Ala Ser Thr Pro Leu Gly Val Gly Gly Phe Gly Ala Ala Arg Ala 195 200 205		
Leu Ser Thr Arg Asn Asp Asn Pro Gln Ala Ala Ser Arg Pro Trp Asp 210 215 220		
Lys Glu Arg Asp Gly Phe Val Leu Gly Asp Gly Ala Gly Met Leu Val 225 230 235 240		
Leu Glu Glu Tyr Glu His Ala Lys Lys Arg Gly Ala Lys Ile Tyr Ala 245 250 255		
Glu Leu Val Gly Phe Gly Met Ser Ser Asp Ala Tyr His Met Thr Ser 260 265 270		
Pro Pro Glu Asn Gly Ala Gly Ala Ala Leu Ala Met Ala Asn Ala Leu 275 280 285		
Arg Asp Ala Gly Ile Glu Ala Ser Gln Ile Gly Tyr Val Asn Ala His 290 295 300		
Gly Thr Ser Thr Pro Ala Gly Asp Lys Ala Glu Ala Gln Ala Val Lys 305 310 315 320		
Thr Ile Phe Gly Glu Ala Ala Ser Arg Val Leu Val Ser Ser Thr Lys 325 330 335		
Ser Met Thr Gly His Leu Leu Gly Ala Ala Gly Ala Val Glu Ser Ile 340 345 350		
Tyr Ser Ile Leu Ala Leu Arg Asp Gln Ala Val Pro Pro Thr Ile Asn 355 360 365		
Leu Asp Asn Pro Asp Glu Gly Cys Asp Leu Asp Phe Val Pro His Glu 370 375 380		
Ala Arg Gln Val Ser Gly Met Glu Tyr Thr Leu Cys Asn Ser Phe Gly 385 390 395 400		
Phe Gly Gly Thr Asn Gly Ser Leu Ile Phe 405 410		

<210> 39

<211> 406
 <212> PRT
 <213> Escherichia coli

<400> 39

Met	Lys	Arg	Ala	Val	Ile	Thr	Gly	Leu	Gly	Ile	Val	Ser	Ser	Ile	Gly
1				5					10					15	
Asn	Asn	Gln	Gln	Glu	Val	Leu	Ala	Ser	Leu	Arg	Glu	Gly	Arg	Ser	Gly
			20					25					30		
Ile	Thr	Phe	Ser	Gln	Glu	Leu	Lys	Asp	Ser	Gly	Met	Arg	Ser	His	Val
		35					40					45			
Trp	Gly	Asn	Val	Lys	Leu	Asp	Thr	Thr	Gly	Leu	Ile	Asp	Arg	Lys	Val
	50					55					60				
Val	Arg	Phe	Met	Ser	Asp	Ala	Ser	Ile	Tyr	Ala	Phe	Leu	Ser	Met	Glu
65					70					75					80
Gln	Ala	Ile	Ala	Asp	Ala	Gly	Leu	Ser	Pro	Glu	Ala	Tyr	Gln	Asn	Asn
				85					90					95	
Pro	Arg	Val	Gly	Leu	Ile	Ala	Gly	Ser	Gly	Gly	Gly	Ser	Pro	Arg	Phe
			100					105						110	
Gln	Val	Phe	Gly	Ala	Asp	Ala	Met	Arg	Gly	Pro	Arg	Gly	Leu	Lys	Ala
		115					120					125			
Val	Gly	Pro	Tyr	Val	Val	Thr	Lys	Ala	Met	Ala	Ser	Gly	Val	Ser	Ala
	130						135					140			
Cys	Leu	Ala	Thr	Pro	Phe	Lys	Ile	His	Gly	Val	Asn	Tyr	Ser	Ile	Ser
145					150					155					160
Ser	Ala	Cys	Ala	Thr	Ser	Ala	His	Cys	Ile	Gly	Asn	Ala	Val	Glu	Gln
				165					170					175	
Ile	Gln	Leu	Gly	Lys	Gln	Asp	Ile	Val	Phe	Ala	Gly	Gly	Gly	Glu	Glu
		180						185						190	
Leu	Cys	Trp	Glu	Met	Ala	Cys	Glu	Phe	Asp	Ala	Met	Gly	Ala	Leu	Ser
		195					200					205			
Thr	Lys	Tyr	Asn	Asp	Thr	Pro	Glu	Lys	Ala	Ser	Arg	Thr	Tyr	Asp	Ala
	210					215					220				
His	Arg	Asp	Gly	Phe	Val	Ile	Ala	Gly	Gly	Gly	Gly	Met	Val	Val	Val
225					230					235					240
Glu	Glu	Leu	Glu	His	Ala	Leu	Ala	Arg	Gly	Ala	His	Ile	Tyr	Ala	Glu
				245					250					255	
Ile	Val	Gly	Tyr	Gly	Ala	Thr	Ser	Asp	Gly	Ala	Asp	Met	Val	Ala	Pro
		260						265					270		

Ser Gly Glu Gly Ala Val Arg Cys Met Lys Met Ala Met His Gly Val
 275 280 285

Asp Thr Pro Ile Asp Tyr Leu Asn Ser His Gly Thr Ser Thr Pro Val
 290 295 300

Gly Asp Val Lys Glu Leu Ala Ala Ile Arg Glu Val Phe Gly Asp Lys
 305 310 315 320

Ser Pro Ala Ile Ser Ala Thr Lys Ala Met Thr Gly His Ser Leu Gly
 325 330 335

Ala Ala Gly Val Gln Glu Ala Ile Tyr Ser Leu Leu Met Leu Glu His
 340 345 350

Gly Phe Ile Ala Pro Ser Ile Asn Ile Glu Glu Leu Asp Glu Gln Ala
 355 360 365

Ala Gly Leu Asn Ile Val Thr Glu Thr Thr Asp Arg Glu Leu Thr Thr
 370 375 380

Val Met Ser Asn Ser Phe Gly Phe Gly Gly Thr Asn Ala Thr Leu Val
 385 390 395 400

Met Arg Lys Leu Lys Asp
 405

<210> 40
 <211> 416
 <212> PRT
 <213> Mycobacterium tuberculosis

<400> 40

Met Ser Gln Pro Ser Thr Ala Asn Gly Gly Phe Pro Ser Val Val Val
 1 5 10 15

Thr Ala Val Thr Ala Thr Thr Ser Ile Ser Pro Asp Ile Glu Ser Thr
 20 25 30

Trp Lys Gly Leu Leu Ala Gly Glu Ser Gly Ile His Ala Leu Glu Asp
 35 40 45

Glu Phe Val Thr Lys Trp Asp Leu Ala Val Lys Ile Gly Gly His Leu
 50 55 60

Lys Asp Pro Val Asp Ser His Met Gly Arg Leu Asp Met Arg Arg Met
 65 70 75 80

Ser Tyr Val Gln Arg Met Gly Lys Leu Leu Gly Gly Gln Leu Trp Glu
 85 90 95

Ser Ala Gly Ser Pro Glu Val Asp Pro Asp Arg Phe Ala Val Val Val
 100 105 110

Gly Thr Gly Leu Gly Gly Ala Glu Arg Ile Val Glu Ser Tyr Asp Leu
 115 120 125

Met	Asn	Ala	Gly	Gly	Pro	Arg	Lys	Val	Ser	Pro	Leu	Ala	Val	Gln	Met	130	135	140	
Ile	Met	Pro	Asn	Gly	Ala	Ala	Ala	Val	Ile	Gly	Leu	Gln	Leu	Gly	Ala	145	150	155	160
Arg	Ala	Gly	Val	Met	Thr	Pro	Val	Ser	Ala	Cys	Ser	Ser	Gly	Ser	Glu	165	170	175	
Ala	Ile	Ala	His	Ala	Trp	Arg	Gln	Ile	Val	Met	Gly	Asp	Ala	Asp	Val	180	185	190	
Ala	Val	Cys	Gly	Gly	Val	Glu	Gly	Pro	Ile	Glu	Ala	Leu	Pro	Ile	Ala	195	200	205	
Ala	Phe	Ser	Met	Met	Arg	Ala	Met	Ser	Thr	Arg	Asn	Asp	Glu	Pro	Glu	210	215	220	
Arg	Ala	Ser	Arg	Pro	Phe	Asp	Lys	Asp	Arg	Asp	Gly	Phe	Val	Phe	Gly	225	230	235	240
Glu	Ala	Gly	Ala	Leu	Met	Leu	Ile	Glu	Thr	Glu	Glu	His	Ala	Lys	Ala	245	250	255	
Arg	Gly	Ala	Lys	Pro	Leu	Ala	Arg	Leu	Leu	Gly	Ala	Gly	Ile	Thr	Ser	260	265	270	
Asp	Ala	Phe	His	Met	Val	Ala	Pro	Ala	Ala	Asp	Gly	Val	Arg	Ala	Gly	275	280	285	
Arg	Ala	Met	Thr	Arg	Ser	Leu	Glu	Leu	Ala	Gly	Leu	Ser	Pro	Ala	Asp	290	295	300	
Ile	Asp	His	Val	Asn	Ala	His	Gly	Thr	Ala	Thr	Pro	Ile	Gly	Asp	Ala	305	310	315	320
Ala	Glu	Ala	Asn	Ala	Ile	Arg	Val	Ala	Gly	Cys	Asp	Gln	Ala	Ala	Val	325	330	335	
Tyr	Ala	Pro	Lys	Ser	Ala	Leu	Gly	His	Ser	Ile	Gly	Ala	Val	Gly	Ala	340	345	350	
Leu	Glu	Ser	Val	Leu	Thr	Val	Leu	Thr	Leu	Arg	Asp	Gly	Val	Ile	Pro	355	360	365	
Pro	Thr	Leu	Asn	Tyr	Glu	Thr	Pro	Asp	Pro	Glu	Ile	Asp	Leu	Asp	Val	370	375	380	
Val	Ala	Gly	Glu	Pro	Arg	Tyr	Gly	Asp	Tyr	Arg	Tyr	Ala	Val	Asn	Asn	385	390	395	400
Ser	Phe	Gly	Phe	Gly	Gly	His	Asn	Val	Ala	Leu	Ala	Phe	Gly	Arg	Tyr	405	410	415	

<210> 41

<211> 438

<212> PRT

<213> Mycobacterium tuberculosis

<400> 41

Met	Gly	Val	Pro	Pro	Leu	Ala	Gly	Ala	Ser	Arg	Thr	Asp	Met	Glu	Gly	
1				5					10					15		
Thr	Phe	Ala	Arg	Pro	Met	Thr	Glu	Leu	Val	Thr	Gly	Lys	Ala	Phe	Pro	
			20					25					30			
Tyr	Val	Val	Val	Thr	Gly	Ile	Ala	Met	Thr	Thr	Ala	Leu	Ala	Thr	Asp	
		35					40					45				
Ala	Glu	Thr	Thr	Trp	Lys	Leu	Leu	Leu	Asp	Arg	Gln	Ser	Gly	Ile	Arg	
	50					55					60					
Thr	Leu	Asp	Asp	Pro	Phe	Val	Glu	Glu	Phe	Asp	Leu	Pro	Val	Arg	Ile	
65					70					75					80	
Gly	Gly	His	Leu	Leu	Glu	Glu	Phe	Asp	His	Gln	Leu	Thr	Arg	Ile	Glu	
			85						90					95		
Leu	Arg	Arg	Met	Gly	Tyr	Leu	Gln	Arg	Met	Ser	Thr	Val	Leu	Ser	Arg	
			100					105					110			
Arg	Leu	Trp	Glu	Asn	Ala	Gly	Ser	Pro	Glu	Val	Asp	Thr	Asn	Arg	Leu	
		115					120						125			
Met	Val	Ser	Ile	Gly	Thr	Gly	Leu	Gly	Ser	Ala	Glu	Glu	Leu	Val	Phe	
	130					135					140					
Ser	Tyr	Asp	Asp	Met	Arg	Ala	Arg	Gly	Met	Lys	Ala	Val	Ser	Pro	Leu	
145					150					155					160	
Thr	Val	Gln	Lys	Tyr	Met	Pro	Asn	Gly	Ala	Ala	Ala	Ala	Val	Gly	Leu	
			165						170					175		
Glu	Arg	His	Ala	Lys	Ala	Gly	Val	Met	Thr	Pro	Val	Ser	Ala	Cys	Ala	
			180					185					190			
Ser	Gly	Ala	Glu	Ala	Ile	Ala	Arg	Ala	Trp	Gln	Gln	Ile	Val	Leu	Gly	
	195						200					205				
Glu	Ala	Asp	Ala	Ala	Ile	Cys	Gly	Gly	Val	Glu	Thr	Arg	Ile	Glu	Ala	
	210					215					220					
Val	Pro	Ile	Ala	Gly	Phe	Ala	Gln	Met	Arg	Ile	Val	Met	Ser	Thr	Asn	
225					230					235					240	
Asn	Asp	Asp	Pro	Ala	Gly	Ala	Cys	Arg	Pro	Phe	Asp	Arg	Asp	Arg	Asp	
			245						250					255		
Gly	Phe	Val	Phe	Gly	Glu	Gly	Gly	Ala	Leu	Leu	Leu	Ile	Glu	Thr	Glu	
		260						265					270			
Glu	His	Ala	Lys	Ala	Arg	Gly	Ala	Asn	Ile	Leu	Ala	Arg	Ile	Met	Gly	

275	280	285
Ala Ser Ile Thr Ser Asp Gly Phe His Met Val Ala Pro Asp Pro Asn		
290	295	300
Gly Glu Arg Ala Gly His Ala Ile Thr Arg Ala Ile Gln Leu Ala Gly		
305	310	315 320
Leu Ala Pro Gly Asp Ile Asp His Val Asn Ala His Ala Thr Gly Thr		
	325	330 335
Gln Val Gly Asp Leu Ala Glu Gly Arg Ala Ile Asn Asn Ala Leu Gly		
	340	345 350
Gly Asn Arg Pro Ala Val Tyr Ala Pro Lys Ser Ala Leu Gly His Ser		
	355	360 365
Val Gly Ala Val Gly Ala Val Glu Ser Ile Leu Thr Val Leu Ala Leu		
	370	375 380
Arg Asp Gln Val Ile Pro Pro Thr Leu Asn Leu Val Asn Leu Asp Pro		
	385	390 395 400
Glu Ile Asp Leu Asp Val Val Ala Gly Glu Pro Arg Pro Gly Asn Tyr		
	405	410 415
Arg Tyr Ala Ile Asn Asn Ser Phe Gly Phe Gly Gly His Asn Val Ala		
	420	425 430
Ile Ala Phe Gly Arg Tyr		
	435	

<210> 42
 <211> 418
 <212> PRT
 <213> Rattus norvegicus

<400> 42

Ser Arg Ala Ser Arg Gln Arg Arg Ala Met Glu Glu Val Val Ile Ala	
1	5 10 15
Gly Met Ser Gly Lys Leu Pro Glu Ser Glu Asn Leu Gln Glu Phe Trp	
	20 25 30
Ala Asn Leu Ile Gly Gly Val Asp Met Val Thr Asp Asp Asp Arg Arg	
	35 40 45
Trp Lys Ala Gly Leu Tyr Gly Leu Pro Lys Arg Ser Gly Lys Leu Lys	
	50 55 60
Asp Leu Ser Lys Phe Asp Ala Ser Phe Phe Gly Val His Pro Lys Gln	
	65 70 75 80
Ala His Thr Met Asp Pro Gln Leu Arg Leu Leu Leu Glu Val Ser Tyr	
	85 90 95

Glu	Ala	Ile	Val	Asp	Gly	Gly	Ile	Asn	Pro	Ala	Ser	Leu	Arg	Gly	Thr	100	105	110
Asn	Thr	Gly	Val	Trp	Val	Gly	Val	Ser	Gly	Ser	Glu	Ala	Ser	Glu	Ala	115	120	125
Leu	Ser	Arg	Asp	Pro	Glu	Thr	Leu	Leu	Gly	Tyr	Ser	Met	Val	Gly	Cys	130	135	140
Gln	Arg	Ala	Met	Met	Ala	Asn	Arg	Leu	Ser	Phe	Phe	Phe	Asp	Phe	Lys	145	150	155
Gly	Pro	Ser	Ile	Ala	Leu	Asp	Thr	Ala	Cys	Ser	Ser	Ser	Leu	Leu	Ala	165	170	175
Leu	Gln	Asn	Ala	Tyr	Gln	Ala	Ile	Arg	Ser	Gly	Glu	Cys	Pro	Ala	Ala	180	185	190
Ile	Val	Gly	Gly	Ile	Asn	Leu	Leu	Leu	Lys	Pro	Asn	Thr	Ser	Val	Gln	195	200	205
Phe	Met	Lys	Leu	Gly	Met	Leu	Ser	Pro	Asp	Gly	Thr	Cys	Arg	Ser	Phe	210	215	220
Asp	Asp	Ser	Gly	Asn	Gly	Tyr	Cys	Arg	Ala	Glu	Ala	Val	Val	Ala	Val	225	230	235
Leu	Leu	Thr	Lys	Lys	Ser	Leu	Ala	Arg	Arg	Val	Tyr	Ala	Thr	Ile	Leu	245	250	255
Asn	Ala	Gly	Thr	Asn	Thr	Asp	Gly	Cys	Lys	Glu	Gln	Gly	Val	Thr	Phe	260	265	270
Pro	Ser	Gly	Glu	Ala	Gln	Glu	Gln	Leu	Ile	Arg	Ser	Leu	Tyr	Gln	Pro	275	280	285
Gly	Gly	Val	Ala	Pro	Glu	Ser	Leu	Glu	Tyr	Ile	Glu	Ala	His	Gly	Thr	290	295	300
Gly	Thr	Lys	Val	Gly	Asp	Pro	Gln	Glu	Leu	Asn	Gly	Ile	Thr	Arg	Ser	305	310	315
Leu	Cys	Ala	Phe	Arg	Gln	Ser	Pro	Leu	Leu	Ile	Gly	Ser	Thr	Lys	Ser	325	330	335
Asn	Met	Gly	His	Pro	Glu	Pro	Ala	Ser	Gly	Leu	Ala	Ala	Leu	Thr	Lys	340	345	350
Val	Leu	Leu	Ser	Leu	Glu	Asn	Gly	Val	Trp	Ala	Pro	Asn	Leu	His	Phe	355	360	365
His	Asn	Pro	Asn	Pro	Glu	Ile	Pro	Ala	Leu	Leu	Asp	Gly	Arg	Leu	Gln	370	375	380
Val	Val	Asp	Arg	Pro	Leu	Pro	Val	Arg	Gly	Gly	Ile	Val	Gly	Ile	Asn	385	390	395

Ser Phe Gly Phe Gly Gly Ala Asn Val His Val Ile Leu Gln Pro Asn
405 410 415

Ala Ser

<210> 43
<211> 401
<212> PRT
<213> Rhizobium sp. Nodulation Protein E

<400> 43

Met Asp Arg Arg Val Val Ile Thr Gly Ile Gly Gly Leu Cys Gly Leu
1 5 10 15

Gly Thr Asn Ala Ala Ser Ile Trp Lys Glu Met Arg Glu Gly Pro Ser
20 25 30

Ala Ile Ser Pro Ile Ile Thr Thr Asp Leu Tyr Asp Leu Glu Gly Thr
35 40 45

Val Gly Leu Glu Ile Lys Ala Ile Pro Glu His Asp Ile Pro Arg Lys
50 55 60

Gln Leu Val Ser Met Asp Arg Phe Ser Leu Leu Ala Val Ile Ala Ala
65 70 75 80

Thr Glu Ala Met Lys Gln Ala Gly Leu Ser Cys Asp Glu Gln Asn Ala
85 90 95

His Arg Phe Gly Ala Ala Met Gly Leu Gly Gly Pro Gly Trp Asp Thr
100 105 110

Ile Glu Glu Thr Tyr Arg Ser Ile Leu Leu Asp Gly Val Thr Arg Ala
115 120 125

Arg Ile Phe Thr Ala Pro Lys Gly Met Pro Ser Ala Ala Ala Gly His
130 135 140

Val Ser Ile Phe Leu Gly Leu Arg Gly Pro Val Phe Gly Val Thr Ser
145 150 155 160

Ala Cys Ala Ala Gly Asn His Ala Ile Ala Ser Ala Val Asp Gln Ile
165 170 175

Arg Leu Gly Arg Ala Asp Val Met Leu Ala Gly Gly Ser Asp Ala Pro
180 185 190

Leu Thr Trp Gly Val Leu Lys Ser Trp Glu Ala Leu Arg Val Leu Ala
195 200 205

Pro Asp Thr Cys Arg Pro Phe Ser Ala Asp Arg Lys Gly Val Val Leu
210 215 220

Gly Glu Gly Ala Gly Met Ala Val Leu Glu Ser Tyr Glu His Ala Ala
225 230 235 240

Ala Arg Gly Ala Thr Met Leu Ala Glu Val Ala Gly Ile Gly Leu Ser
245 250 255

Gly Asp Ala Tyr Asp Ile Val Met Pro Ser Ile Glu Gly Pro Glu Ala
260 265 270

Ala Met Arg Ser Cys Leu Ala Asp Ala Glu Leu Asn Pro Asp Asp Val
275 280 285

Asp Tyr Leu Asn Ala His Gly Thr Gly Thr Val Ala Asn Asp Glu Met
290 295 300

Glu Thr Ala Ala Ile Lys Arg Val Phe Gly Asp His Ala Phe Gln Met
305 310 315 320

Ser Val Ser Ser Thr Lys Ser Met His Ala His Cys Leu Gly Ala Ala
325 330 335

Ser Ala Leu Glu Met Ile Ala Cys Val Met Ala Ile Gln Glu Gly Val
340 345 350

Ile Pro Pro Thr Ala Asn Tyr Arg Glu Pro Asp Pro Gln Cys Asp Leu
355 360 365

Asp Val Thr Pro Asn Val Pro Arg Glu Gln Arg Cys Gly Ser Met Ser
370 375 380

Asn Ala Phe Ala Met Gly Gly Thr Asn Ala Val Leu Ala Phe Arg Gln
385 390 395 400

Val

<210> 44
<211> 419
<212> PRT
<213> Streptomyces polyketide synthase

<400> 44

Val Asn Arg Arg Ile Val Ile Thr Gly Ile Gly Val Val Ala Pro Gly
1 5 10 15

Ala Val Gly Thr Lys Pro Phe Trp Glu Leu Leu Leu Ser Gly Thr Thr
20 25 30

Ala Thr Arg Ala Ile Ser Thr Phe Asp Ala Thr Pro Phe Arg Ser Arg
35 40 45

Ile Ala Ala Glu Cys Asp Phe Asp Pro Val Ala Ala Gly Leu Ser Ala
50 55 60

Glu Gln Ala Arg Arg Leu Asp Arg Ala Gly Gln Phe Ala Leu Val Ala
65 70 75 80

Gly Gln Glu Ala Leu Ala Asp Ser Gly Leu Arg Ile Asp Glu Asp Ser

85					90					95						
Ala	His	Arg	Val	Gly	Val	Cys	Val	Gly	Thr	Ala	Val	Gly	Cys	Thr	Gln	
100					105					110						
Lys	Leu	Glu	Ser	Glu	Tyr	Val	Ala	Leu	Ser	Ala	Gly	Gly	Ala	His	Trp	
115					120					125						
Val	Val	Asp	Pro	Gly	Arg	Gly	Ser	Pro	Glu	Leu	Tyr	Asp	Tyr	Phe	Val	
130					135					140						
Pro	Ser	Ser	Leu	Ala	Ala	Glu	Val	Ala	Trp	Leu	Ala	Gly	Ala	Glu	Gly	
145					150					155					160	
Pro	Val	Asn	Ile	Val	Ser	Ala	Gly	Cys	Thr	Ser	Gly	Ile	Asp	Ser	Ile	
165					170					175						
Gly	Tyr	Ala	Cys	Glu	Leu	Ile	Arg	Glu	Gly	Thr	Val	Asp	Ala	Met	Val	
180					185					190						
Ala	Gly	Gly	Val	Asp	Ala	Pro	Ile	Ala	Pro	Ile	Thr	Val	Ala	Cys	Phe	
195					200					205						
Asp	Ala	Ile	Arg	Ala	Thr	Ser	Asp	His	Asn	Asp	Thr	Pro	Glu	Thr	Ala	
210					215					220						
Ser	Arg	Pro	Phe	Ser	Arg	Ser	Arg	Asn	Gly	Phe	Val	Leu	Gly	Glu	Gly	
225					230					235					240	
Gly	Ala	Ile	Val	Val	Leu	Glu	Glu	Ala	Glu	Ala	Ala	Val	Arg	Arg	Gly	
245					250					255						
Ala	Arg	Ile	Tyr	Ala	Glu	Ile	Gly	Gly	Tyr	Ala	Ser	Arg	Gly	Asn	Ala	
260					265					270						
Tyr	His	Met	Thr	Gly	Leu	Arg	Ala	Asp	Gly	Ala	Glu	Met	Ala	Ala	Ala	
275					280					285						
Ile	Thr	Ala	Ala	Leu	Asp	Glu	Ala	Arg	Arg	Asp	Pro	Ser	Asp	Val	Asp	
290					295					300						
Tyr	Val	Asn	Ala	His	Gly	Thr	Ala	Thr	Lys	Gln	Asn	Asp	Arg	His	Glu	
305					310					315					320	
Thr	Ser	Ala	Phe	Lys	Arg	Ser	Leu	Gly	Glu	His	Ala	Tyr	Arg	Val	Pro	
325					330					335						
Ile	Ser	Ser	Ile	Lys	Ser	Met	Ile	Gly	His	Ser	Leu	Gly	Ala	Val	Gly	
340					345					350						
Ser	Leu	Glu	Val	Ala	Ala	Thr	Ala	Leu	Ala	Val	Glu	Tyr	Gly	Val	Ile	
355					360					365						
Pro	Pro	Thr	Ala	Asn	Leu	His	Asp	Pro	Asp	Pro	Glu	Leu	Asp	Leu	Asp	
370					375					380						
Tyr	Val	Pro	Leu	Thr	Ala	Arg	Glu	Lys	Arg	Val	Arg	His	Ala	Leu	Thr	

385 390 395 400
 Val Gly Ser Gly Phe Gly Gly Phe Gln Ser Ala Met Leu Leu Ser Arg
 405 410 415

 Leu Glu Arg

 <210> 45
 <211> 416
 <212> PRT
 <213> Synechocystis sp.

 <400> 45

 Met Ala Asn Leu Glu Lys Lys Arg Val Val Val Thr Gly Leu Gly Ala
 1 5 10 15

 Ile Thr Pro Ile Gly Asn Thr Leu Gln Asp Tyr Trp Gln Gly Leu Met
 20 25 30

 Glu Gly Arg Asn Gly Ile Gly Pro Ile Thr Arg Phe Asp Ala Ser Asp
 35 40 45

 Gln Ala Cys Arg Phe Gly Gly Glu Val Lys Asp Phe Asp Ala Thr Gln
 50 55 60

 Phe Leu Asp Arg Lys Glu Ala Lys Arg Met Asp Arg Phe Cys His Phe
 65 70 75 80

 Ala Val Cys Ala Ser Gln Gln Ala Ile Asn Asp Ala Lys Leu Val Ile
 85 90 95

 Asn Glu Leu Asn Ala Asp Glu Ile Gly Val Leu Ile Gly Thr Gly Ile
 100 105 110

 Gly Gly Leu Lys Val Leu Glu Asp Gln Gln Thr Ile Leu Leu Asp Lys
 115 120 125

 Gly Pro Ser Arg Cys Ser Pro Phe Met Ile Pro Met Met Ile Ala Asn
 130 135 140

 Met Ala Ser Gly Leu Thr Ala Ile Asn Leu Gly Ala Lys Gly Pro Asn
 145 150 155 160

 Asn Cys Thr Val Thr Ala Cys Ala Ala Gly Ser Asn Ala Ile Gly Asp
 165 170 175

 Ala Phe Arg Leu Val Gln Asn Gly Tyr Ala Lys Ala Met Ile Cys Gly
 180 185 190

 Gly Thr Glu Ala Ala Ile Thr Pro Leu Ser Tyr Ala Gly Phe Ala Ser
 195 200 205

 Ala Arg Ala Leu Ser Phe Arg Asn Asp Asp Pro Leu His Ala Ser Arg
 210 215 220

Pro Phe Asp Lys Asp Arg Asp Gly Phe Val Met Gly Glu Gly Ser Gly
 225 230 235 240
 Ile Leu Ile Leu Glu Glu Leu Glu Ser Ala Leu Ala Arg Gly Ala Lys
 245 250 255
 Ile Tyr Gly Glu Met Val Gly Tyr Ala Met Thr Cys Asp Ala Tyr His
 260 265 270
 Ile Thr Ala Pro Val Pro Asp Gly Arg Gly Ala Thr Arg Ala Ile Ala
 275 280 285
 Trp Ala Leu Lys Asp Ser Gly Leu Lys Pro Glu Met Val Ser Tyr Ile
 290 295 300
 Asn Ala His Gly Thr Ser Thr Pro Ala Asn Asp Val Thr Glu Thr Arg
 305 310 315 320
 Ala Ile Lys Gln Ala Leu Gly Asn His Ala Tyr Asn Ile Ala Val Ser
 325 330 335
 Ser Thr Lys Ser Met Thr Gly His Leu Leu Gly Gly Ser Gly Gly Ile
 340 345 350
 Glu Ala Val Ala Thr Val Met Ala Ile Ala Glu Asp Lys Val Pro Pro
 355 360 365
 Thr Ile Asn Leu Glu Asn Pro Asp Pro Glu Cys Asp Leu Asp Tyr Val
 370 375 380
 Pro Gly Gln Ser Arg Ala Leu Ile Val Asp Val Ala Leu Ser Asn Ser
 385 390 395 400
 Phe Gly Phe Gly Gly His Asn Val Thr Leu Ala Phe Lys Lys Tyr Gln
 405 410 415

<210> 46
 <211> 441
 <212> PRT
 <213> *Vibrio harveyi*

<400> 46

Ser Asp Tyr His Asn His Phe Ile Asn Val Lys Ala Val Ala Arg Pro
 1 5 10 15
 Leu Phe Phe Cys Leu Phe Trp Arg Thr Ser Val Ala Asn Asn Arg Arg
 20 25 30
 Val Val Ile Thr Gly Leu Gly Ile Val Ser Pro Val Gly Asn Thr Val
 35 40 45
 Ala Thr Ala Trp Glu Ala Ile Lys Ser Gly Ile Ser Gly Ile Glu Asn
 50 55 60
 Ile Glu His Phe Asp Thr Thr Asn Phe Ser Thr Lys Phe Ala Gly Leu
 65 70 75 80

Val	Asn	Asp	Phe	Asp	Ala	Glu	Ser	Val	Gly	Ile	Asn	Arg	Lys	Asp	Cys	85	90	95
Arg	Lys	Met	Asp	Leu	Phe	Ile	Gln	Tyr	Gly	Ile	Ala	Ala	Ala	Glu	Gln	100	105	110
Ala	Leu	Thr	Asp	Ser	Gly	Leu	Glu	Ile	Thr	Glu	Gln	Asn	Ala	Thr	Arg	115	120	125
Ile	Gly	Thr	Ala	Ile	Gly	Ser	Gly	Ile	Gly	Gly	Leu	Gly	Leu	Ile	Glu	130	135	140
Gln	Asn	Val	His	Ser	Phe	Val	Lys	Gly	Gly	Ala	Arg	Lys	Val	Ser	Pro	145	150	155
Phe	Phe	Val	Pro	Ala	Thr	Ile	Val	Asn	Met	Val	Ala	Gly	His	Val	Ser	165	170	175
Ile	Arg	Asn	Asn	Leu	Lys	Gly	Pro	Asn	Ile	Ala	Ile	Ala	Thr	Ala	Cys	180	185	190
Thr	Ser	Gly	Thr	His	Cys	Ile	Gly	Gln	Ser	Ala	Arg	Met	Ile	Ala	Tyr	195	200	205
Gly	Asp	Ala	Asp	Val	Met	Val	Ala	Gly	Gly	Ala	Glu	Lys	Ala	Ser	Thr	210	215	220
Glu	Met	Gly	Leu	Ala	Gly	Phe	Gly	Ser	Ala	Lys	Ala	Leu	Ser	Thr	Arg	225	230	235
Asn	Asp	Asp	Pro	Gln	Lys	Ala	Ser	Arg	Pro	Trp	Asp	Lys	Asp	Arg	Asp	245	250	255
Gly	Phe	Val	Leu	Gly	Asp	Gly	Ala	Gly	Val	Leu	Val	Met	Glu	Glu	Tyr	260	265	270
Glu	His	Ala	Val	Ala	Arg	Gly	Ala	Thr	Ile	Tyr	Ala	Glu	Leu	Ala	Gly	275	280	285
Phe	Gly	Met	Ser	Gly	Asp	Ala	Phe	His	Met	Thr	Ser	Pro	Pro	Glu	Asp	290	295	300
Gly	Ala	Gly	Ala	Ala	Leu	Ser	Met	Asn	Asn	Ala	Ile	Ala	Asp	Ala	Gly	305	310	315
Ile	Thr	Ala	Asp	Lys	Val	Gly	Tyr	Val	Asn	Ala	His	Gly	Thr	Ser	Thr	325	330	335
Pro	Ala	Gly	Asp	Lys	Ala	Glu	Thr	Ala	Ala	Val	Lys	Ser	Val	Phe	Gly	340	345	350
Glu	His	Ala	Tyr	Thr	Leu	Ala	Val	Ser	Ser	Thr	Lys	Ser	Met	Thr	Gly	355	360	365
His	Leu	Leu	Gly	Ala	Ala	Gly	Ala	Ile	Glu	Ala	Ile	Phe	Thr	Ile	Leu	370	375	380

Ala Leu Lys Asp Gln Ile Leu Pro Pro Thr Ile Asn Leu Glu Asn Pro
385 390 395 400

Ser Glu Gly Cys Asp Leu Asp Tyr Val Thr Asp Gly Ala Arg Pro Val
405 410 415

Asn Met Glu Tyr Ala Leu Ser Asn Ser Phe Gly Phe Gly Gly Thr Asn
420 425 430

Gly Ser Leu Leu Phe Lys Lys Ala Asp
435 440